

INTENSE PULSED NEUTRON SOURCE



POSY1 *A Users Guide*

OCTOBER 2004

ARGONNE NATIONAL LABORATORY
INTENSE PULSED NEUTRON SOURCE
9700 SOUTH CASS AVENUE
ARGONNE, ILLINOIS 60439
PHONE: (630) 252-4328
FAX: (630) 252-4163
E-MAIL: RJGOYETTE@ANL.GOV



Disclaimer

©1993, 1994, 1995, 1996, 1997, 1998, 1999, 2003, 2004 Argonne National Laboratory, Intense Pulsed Neutron Source

This procedure was prepared for the purpose of assisting visitors who are performing work sponsored by an agency of the United States Government. Neither the United States Government nor any agency thereof, nor any of their employees, makes any warranty, express or implied, or assumes any legal liability or responsibility for the accuracy, completeness, or usefulness of any information, apparatus, product, or process disclosed, or represents that its use would not infringe privately owned rights. Reference herein to any specific commercial product, process, or service by trade name, trademark, manufacturer, or otherwise does not necessarily constitute or imply its endorsement, recommendation, or favoring by the United States Government or any agency thereof. The views and opinions of the authors expressed herein do not necessarily state or reflect those of the United States Government or any agency thereof.



Argonne National Laboratory
Intense Pulsed Neutron Source
9700 South Cass Avenue
Argonne, Illinois 60439
(630)252-6485



Acknowledgements

The current data analysis, fitting and plotting package, called POSYIDL2, is based on software originally written for IPNS using Interactive Data Language (IDL) from Research Systems, Inc. by Dr. Alan Munter.



Contact Information

Rick Goyette, Office:	2-4328
Pager:	266-8329
Home:	778-1324

Suzanne te Velthuis, Office:	2-1075
Home:	(773) 404-1388
Cell:	(773) 515-2296

Abdel Alsmadi, Office:	2-3775
Cell:	(505) 571-3490

Accelerator Main Control Room:	2-6562
--------------------------------	--------

Health Physics Technician

Mark Delrose, Office:	2-8185
Pager:	266-0909

Randy Wilson, Office:	2-7170
-----------------------	--------

After normal buisness hours	471-8792
-----------------------------	----------

ALL EMERGENCIES	911
------------------------	------------



IPNS Operations

While the IPNS facility operates seven days a week, twenty four hours a day during normal operations, there are scheduled periods during operations cycles when IPNS does not produce neutrons for users. These periods are as follows:

RCS maintenance checks:

Daily 9:00 A.M. - 9:15 A.M.

Accelerator machine research periods:

First Thursday of every run cycle 9:00 A.M. - 11:00 A.M.

Every Tuesday 9:00 A.M. - 3:00 P.M.

Moderator annealing:

Every Monday and Friday, 9:00 A.M. - 9:45 A.M.

Moderator gas exchange:

Every Tuesday, during machine research, 9:00 A.M. - 3:00 P.M.

No data will be collected during these periods. This is normal.



Table of Contents

Opening the PNE Directory.....	12
Defining a New Run.....	12
Changing Control Parameters.....	14
Changing Sample Angle	16
Changing Analyzer Position.....	16
Changing Sample Magnetic Field.....	17
Changing the Sample Temperature.....	17
Translating the Sample Stage.....	19
Checking Control Parameters.....	19
Starting and Stopping a Run.....	20
Scheduling Runs	21
Checking the Status of a Live Run.....	22
Modifying Control Parameters.....	22
Reading TO Pulses Using RDP.....	23
Reading Sample Temperature.....	23
Intensity as a Function of Position.....	24
Intensity as a Function of Time-of-Flight.....	25
Rescaling the X and Y Axis of Any Plot.....	25
Changing the Y-Axis.....	25
Changing the X-Axis.....	26
Looking at Different Spin States.....	26
Integrating Over Several Channels.....	27
Looking at Old Data.....	27
Looking at Monitor Data.....	27
Reinitializing the Graphics Software.....	27
Operator Messages.....	28
Setting Up a Transmission Run.....	29
Processing a Transmission Run.....	29
Plotting a Transmission Data File.....	31
Generating a Reflectivity File.....	33
Creating a .DAT File Using Spinread.....	33

Createing a Reflectivity File Using SPINREF.....	34
Calculating Polarization using POL.....	35
Merging Reflectivity Files.....	36
Plotting Reflectivity Files.....	37
Plotting Contours of Run Files.....	38
Fitting Data Using LAYERS.....	40
Performing a Full System Reboot.....	42
Performing a Minimal System Reboot.....	42
Instructions on using the POSY1 analyzer.....	44
Introduction to Reflection Theory	47
Reflection Theory	48
Instrumentation	55
Isotopic Substitution	58
Data Analysis	59
Model Fitting	60
Maximum Entropy and Bayesian Analysis	62
Speckle Holography and Other Techniques	63
Inversion Techniques	66
References	67



Setting Up and Starting Run Files

This Chapter describes how to set up and start a new run, or a sequence of new runs, stop an active run and schedule a run, or a sequence of runs. It is assumed that all run file set up will be done from the DEC Windows terminal (the big screen by the computer). In all instances, <cr> means carriage return.

NOTE: Pay particular attention to the setting up process: it contains several parts and is the most important part of the data collection process. Discuss the set up parameters beforehand, and decide on the final settings before you start setting up a run file.

Opening the PNE Directory

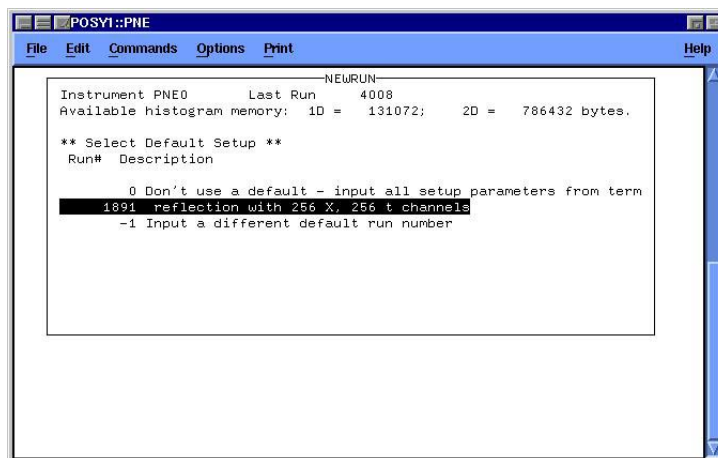
To move to the PNE directory, type SET DEF [PNE] <cr> (make sure there is a space between DEF and [PNE]). All run files are stored in [PNE.DATA].

Defining a New Run

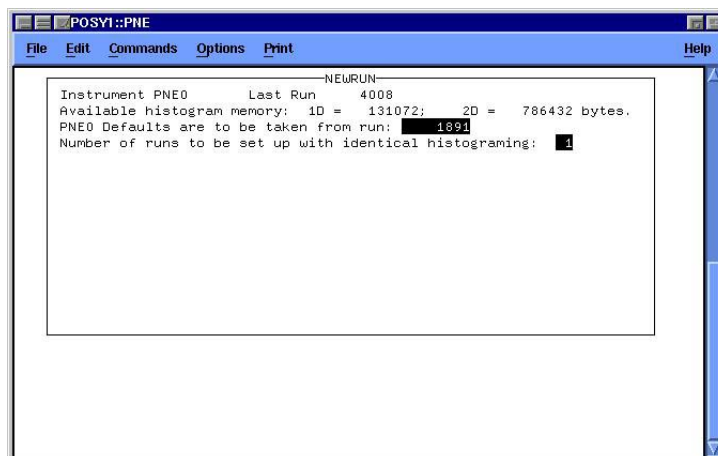
To define a new run, use the following procedure:

Type NEWRUN <cr>.

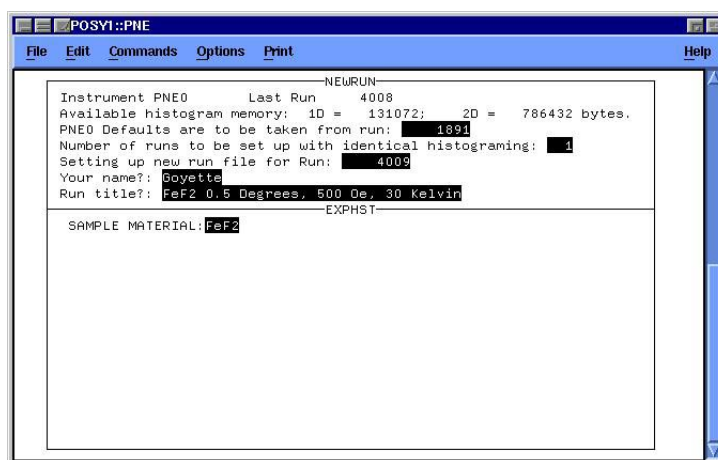
The MicroVax allows you to adjust the settings of the run by prompting you with a list of questions. Some of these settings have default values which will appear after the prompt in a black highlighted region. These should normally be accepted, by pressing the carriage return (<cr>). First, you will have to choose a run file from which to read the parameters for this run file:



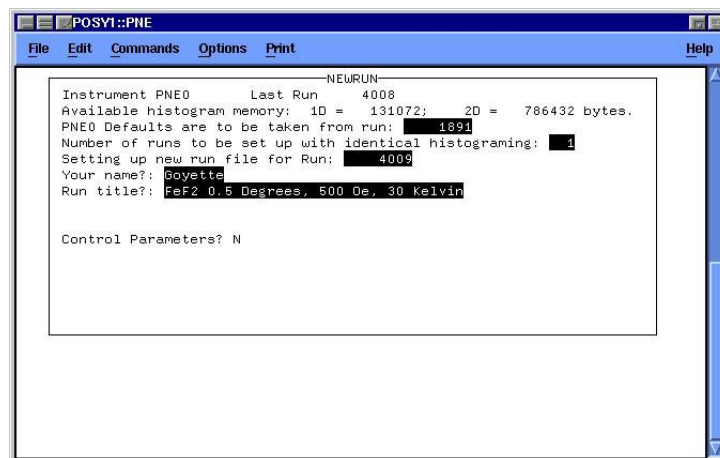
Run number 1891 is the current default run that we use to set up a run file with 256 position channels and 256 time channels. Do not change the default run number without first consulting with a member of the POSY staff.



You will also be asked how many runs to set up with identical histogramming. This number will usually be one. Consult a member of the POSY staff for setting up more than one run at a time. Next you will be asked to supply some information on the new run:



After which you will be placed back in the main menu. Next you will have the option of changing some control parameters:



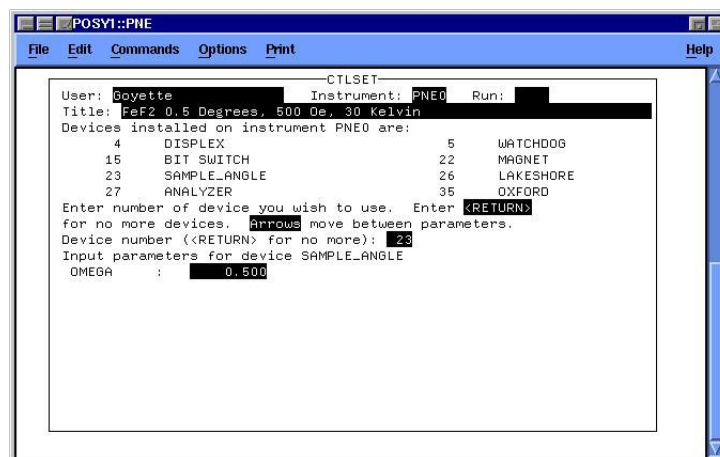
If you do not want to change any of the control parameters (for example, the angle) then enter a carriage return to complete the set up of the run.

Changing Control Parameters

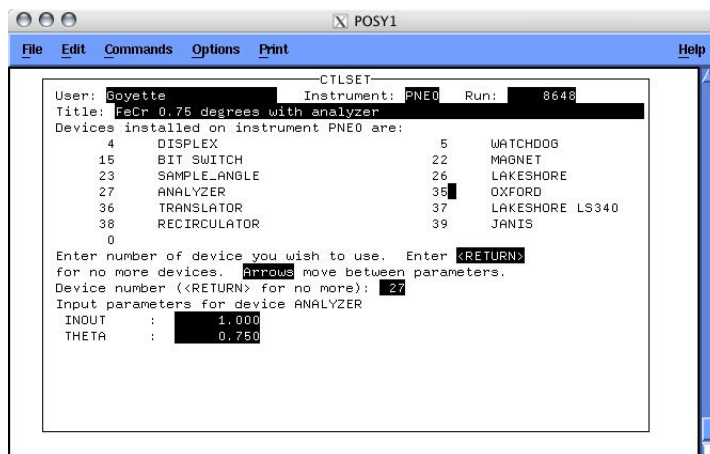
If you wish to change some control parameters, you should still be at the prompt :

Control Parameters: {N}.

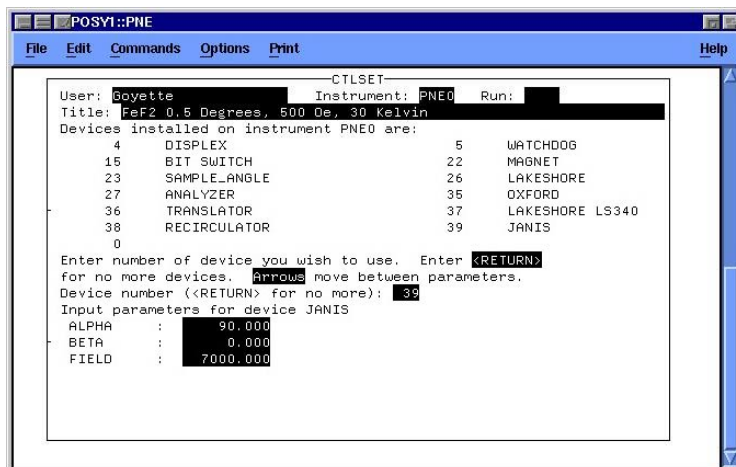
Type: Y <cr> . This will bring up an additional menu. Choosing the default setting of zero, or typing zero if the device is set to another number, will put you back to the Newrun menu. Currently, Sample_angle, Analyzer, Janis and Lakeshore LS340 are the only devices available for use. To change the angle of the sample, choose Device Number 23:



Input the desired sample angle in degrees. When the run is started the computer will automatically set the sample angle. To use the Analyzer, choose Device Number 27. Two parameters will be required: INOUT and THETA. When controlled by a run file, any positive number entered for INOUT will cause the analyzer to be placed in the beam, and the angle entered in THETA will be used in determining the analyzer position. When controlled from a run file, and negative number entered for INOUT will cause the analyzer to be removed from the beam and rotated to a flat angle. The number entered in THETA will not be used.

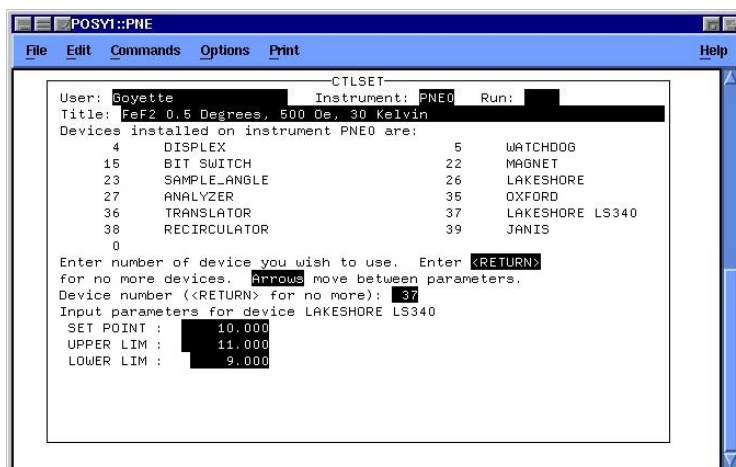


To use the electromagnet, choose Device Number 22. Enter the sample magnetic field in Oersted (gauss).



Ask a member of the POSY1 staff, or refer to the diagram posted by the instrument for information on how angles alpha and beta are defined.

If you are using the JANIS Cryostat, you may wish to set the sample temperature from the run file. To set the sample temperature, use device number 37 (Lakeshore LS340):



Enter the desired set point at the prompt. This is the temperature the Lakeshore controller will hold. Next, enter an upper and a lower limit for the temperature. This is a temperature window that is acceptable to you for data collection. If the temperature moves out of this window, the runtime software will inhibit data collection until the temperature moves back into this window.

Changing Sample Angle

The program ANGLE is provided on the POSY1 system. It is used to change the sample angle. To use this program, type ANGLE <cr>. You will be prompted:

```
ENTER NEW SAMPLE ANGLE (P), CHANGE LIMITS (L),  
CALIBRATE (C), CHANGE OTHER PARAMETERS (O),  
OR EXIT (E) ?
```

Type P to change the sample angle. You will be prompted again:

```
USE <CR> FOR NO CHANGE  
CURRENT ANGLE=  0.90  DEGREES.  NEW ANGLE ?
```

Type in the new angle and hit return to set the new angle, or hit return to exit without changing the angle. The computer will accept any angle from -20 to + 20 degrees. . It will take a moment to drive the sample to the new angle.

Changing Analyzer Position

Use the program ANALYZER to insert or remove the analyzer interactively by typing ANALYZER <cr> at the command prompt:

```
POSY1_PNE> analyzer  
*** ANALYZER ***  
PNE0 USING ANALYZER          -- DEVICE    27  
MIRROR POSITION                OUT  
MIRROR ANGLE (THETA)          -0.300  
THETA0                        0.600  
CALCULATED ANGLE (PHI)        0.000
```

```
CHANGE MIRROR POSITIONING (AUTOMATIC TRANSLATION AND ROTATION) (P)  
CALIBRATE CURRENT MIRROR ANGLE (C)  
CHANGE OTHER PARAMETERS (THETA0, X0, IN POS OR OUT POS) (O)  
PERFORM DIAGNOSTICS (TRANSLATE OR ROTATE MANUALLY) (D)  
CHANGE LIMITS (L)  
READ CURRENT TRANSLATION POSITION (R)  
EXIT (E)  
ENTER CHOICE: p
```

Type P to change the analyzer position. When moving the analyzer IN you will be asked to specify a mirror angle. When moving the analyzer OUT you will not be asked for a mirror angle.

```
MIRROR POSITION IS CURRENTLY OUT.  NEW POSITION (IN OR OUT): IN  
MIRROR ROTATION (THETA) IS CURRENTLY -0.30 DEGREES.  NEW ROTA-  
TION:0.75
```

Mirror translations will be calculated automatically from desired mirror angles when option P is selected. In the event that manual translation of the mirror is desired, using

option D, note that entering a destination position more positive than the current position will translate the analyzer to the right (from a viewers perspective if you are standing at the instrument, facing the cryostat, with your back to the detector), while a destination position more negative than the current position will translate the analyzer to the left. Also note that this behavior is opposite the behavior for translating the sample stage.

Changing Sample Magnetic Field

Use the program JANIS to manually change the magnetic field at the sample position by typing JANIS <cr>. you will be prompted:

```
PNE0 USING JANIS          -- DEVICE    39

HELIUM LEVEL =  75.00%.  NITROGEN LEVEL =  89.00%.
```

```
>>>>>>>>> ALL COILS ARE AT ZERO <<<<<<<<<<
```

```
CURRENT ANGLE ALPHA IS    0.00 DEGREES.
CURRENT ANGLE BETA  IS    0.00 DEGREES.
CURRENT FIELD STRENGTH IS      0.00 GAUSS.
FIELD STEP SIZE (OE):      0.
RAMP/ROTATE IS SET TO RAMP.
X-COIL FIELD (OE):        0.
Y-COIL FIELD (OE):        0.
Z-COIL FIELD (OE):        0.
```

```
ENTER NEW FIELD SETTINGS (P), CHANGE LIMITS (L),
OR EXIT (E)? (DEFAULT=EXIT):
```

Type P to change the field strength or E to exit without making any changes. You will be prompted again:

```
ANGLE ALPHA IS    0.00 DEGREES.  NEW ALPHA (IN DEGREES)?: 90.
ALGLE BETA  IS    0.00 DEGREES.  NEW BETA  (IN DEGREES)?: 0.
FIELD STRENGTH IS      0.00 GAUSS. NEW FIELD  (IN GAUSS)?: 7999.
FIELD STEP SIZE IS      0.000 GAUSS. NEW STEP SIZE (IN GAUSS,
zero=no limit)?:0
Maximum step size will be used
Ramp (0) or Rotate (1) (Default is RAMP): 0
```

Ask a member of the POSY1 ataff, or refer to the diagram posted by the instrument for information on how angles alpha and beta are defined.

Changing the Sample Temperature

The program TEMTUR will allow you to change the sample temperature. To use it, type TEMTUR <cr>. You must specify which device to use. The normal device installed on

POSY1 is device 37 (Lakeshore LS340). Specify device 37 at the prompt:

DEVICES AVAILABLE ARE :

4	DISPLEX
26	LAKESHORE
35	OXFORD
37	LAKESHORE LS340
38	RECIRCULATOR

ENTER ----> DEVICE NUMBER: 37 <cr>

Now you will have the choice of entering the setpoint or changing other parameters.

S = ENTER SETPOINT AND LIMITS ONLY - DOWNLOAD ALL PARAMETERS TO THE LAKESHORE CONTROLLER
C = CHANGE OTHER CONTROLLER PARAMETERS - NO DOWNLOADING OF PARAMETERS - USE "S" ABOVE TO DOWNLOAD ALL PARAMETERS TO THE LAKESHORE CONTROLLER
E = EXIT

ENTER --> S, C or E:

You should not attempt to change any other parameters without first consulting with a member of the POSY1 staff. Type S at the prompt to change the sample temperature. You will get the set point and limits prompt. Enter the desired sample temperature and limits. The default values are in parenthesis. Do not enter a temperature above 300 K.

SET POINT ?	(50.0)100.0
UPPER LIMIT ?	(55.0)105.0
LOWER LIMIT ?	(45.0)95.0

The program will show you the settings after you have entered them.

SET POINT	100.000	UPPER LIM	105.000	LOWER LIM	
95.000 DELAY	1.000				
CTL SENS	1.000	MEAS SENS	1.000	TUNE	2.000
GAIN	6.000				
RESET	9.000	RATE	0.100	HEATER	3.000 RAMP 0-
1	0.000				
RAMP RATE	0.000				

Make sure they are correct. The program will display a warning concerning the possible effects of downloading an incorrect temperature to the controller. If the values are not correct type E to exit and re-run TEMTUR to enter the correct values. If the values are correct type DOWNLOAD at the prompt.

E = Exit TEMTUR without downloading
DOWNLOAD = Continue and download control-
ler parameters.

ENTER --> E or DOWNLOAD :

Translating the Sample Stage

The sample stage can be translated perpendicular to the plane of the incident beam by using the command TRANSLATE and selecting option P:

```
POSY1_PNE> translate
*** TRANSLATE ***
PNE0 USING TRANSLATOR      -- DEVICE    36

ENTER NEW SAMPLE POSITION (P), READ CURRENT SAMPLE PO-
SITION (R),
CHANGE LIMITS (L), CHANGE OTHER PARAMETERS (O),
OR EXIT (E) ?
P
USE <CR> FOR NO CHANGE
CURRENT POSITION=  -0.41  MILLIMETERS.  NEW POSITION ?
```

Enter the desired destination position in millimeters. Note that entering a destination position more positive than the current position will translate the sample to the left (from a viewers perspective if you are standing at the instrument, facing the cryostat, with your back to the detector), while a destination position more negative than the current position will translate the sample to the right. Also note that this behavior is opposite the behavior for translating the analyzer.

Checking Control Parameters

Before typing START, it would be wise to check the control parameters by using the command PRUN CONTROL RUN #.

When you use the Prun Control command, the computer will display all control parameters that were set for that run number. If you set the angle as a control parameter, the current angle and the angle that will be set for the next run will be displayed:

```

POSY1_PNE> prun control 4009
N32016 MICROPROCESSOR - 1 PROCESSORS

PNE0 RUN      4009: Goyette
FeF2 0.5 Degrees, 500 Oe, 30 Kelvin

Controlled Devices and Parameter Settings
DEVICE: SAMPLE_ANGLE
OMEGA      0.500 OMEGA NOW      0.900 OMEGA LOW  -20.000 OMEGA HI   20.000
OMEGA STP   4.000 OMEGA BK      100.000 OMEGA MOT   1.000
DEVICE: MAGNET
Ew FIELD    500.000 FIELD NOW      0.000 HICAL      3231.100 LOCAL      310.840
DEVICE: OXFORD
SET POINT   30.000 UPPER LIM      32.000 LOWER LIM   28.000 DELAY      5.000
CTL SENS     1.000 MEAS SENS       1.000 HEATA/M      3.000 PID A/M      1.000
GAIN         12.000 RESET          20.000 RATE         50.000 GAS (M)     24.000
HEATER       0.000 MAX HEAT        0.000
POSY1_PNE>
  
```

NOTE: PRUN CONTROL will only give information on the angle, magnetic field and temperature if they were set as control parameters. It will not give any information if the parameters were set using the programs ANGLE, MAGNET or OXFORD.

Starting and Stopping a Run

To start the new run type START run # <cr>. You will be asked to verify scheduling and setup of the run:

```

START
Starting runs for Instrument : PNE0
The following runs are scheduled:
PNE0 run      4009 set up by Goyette
The Runs are in the following sequence:
4009
4009 has      720 cycles of 108000 pulses.

Is this the sequence and monitoring you want? Y
  
```

Type Y at the prompt if the information is correct to start the run. Type N to re-enter the start command with the proper information.

If you wish to stop the run and save the data currently in the multibus memory, use the command:

STOPRUN <cr>.

To stop the run and not save the data in the multibus memory, use the command: QUITRUN <cr>. If you used QUITRUN by accident where you wanted to use STOPRUN, just type: START/RESTART RUN #, which will pick up where you left off when you quit, then use STOPRUN, which will save the data.

START/RESTART will also allow you to resume data collection after system malfunctions, without losing the data currently in the multibus memory. If you have stopped the run accidentally using the STOPRUN command, type START run#, where run# is the run you accidentally stopped. The computer will ask you if you wish to erase the data in the multibus memory. Type N for NO and the computer will start collecting data at the point where it left off when you typed stop.

Normally, the data are stored in the multibus memory and every hour during an active run are written to the directory [PNE.DATA]. The computer will keep the latest three versions of the data set automatically. Every time the multibus memory is dumped to [PNE.DATA] the run file version will be updated.

Remember:

STOPRUN—saves data QUITRUN—does not save data

Scheduling Runs

If you want to schedule a run to run for a set amount of time, use the schedule command:

```
SCHEDULE TIME(RUN#)/# OF PULSES <cr>
```

There are 108000 Pulses in one hour, so if you want a run to last one hour then use 108000P for # of pulses, and the number 1 for time. If you want a run to be 30 min., then use 54000P for # of pulses, and 1 for time. Lets say that you wanted to schedule three runs in a row, and that these runs have run #'s 500,501 and 502. Lets also assume that you want the first run to last two hour, the second run to last one-half hour, and the third run to last three-quarters of an hour. Your first schedule command would look like this:

```
SCHEDULE 2(500)/108000P <cr>
```

the second schedule command would look like:

```
SCHEDULE 1(501)/54000P <cr>
```

and the third schedule command would be:

```
SCHEDULE 3(502)/27000P <cr>.
```

Notice that for the third command, we have specified the number of pulses corresponding to one-quarter of an hour, but we have defined the time of the run to be three of these increments: three groups of pulses at 1/4 hour per group is 3/4 hour for run 502. You must set the parameters for each run using the NEWRUN command, before you can schedule the run. If you have not done this before, refer to the section on STARTING A NEW RUN. You may check the scheduling of a run using PRUN CONTROL. If you have scheduled more than one run and you want to check them all then use the same command, but list all the runs you want to check:

```
PRUN CONTROL 500,501,502 <cr>.
```

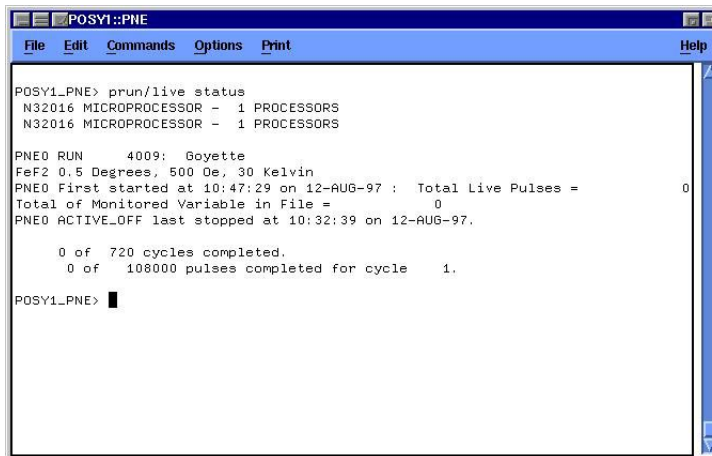
You may use the START command to start more than run file. For example, to start the example runs above, you would type:

START 500,501,502 <cr>.

This command will start run 500 and run it for the scheduled amount of time. once run 500 is finished, 501 will start and run for its scheduled amount of time. Finally, run 502 will be started and will run for its scheduled time.

Checking the Status of a Live Run

To determine whether a live run is active, or to determine the status of a run that is known to be live, use the command PRUN/LIVE STATUS. The computer will display information concerning the live run:



```
POSY1_PNE> prun/live status
N32016 MICROPROCESSOR - 1 PROCESSORS
N32016 MICROPROCESSOR - 1 PROCESSORS

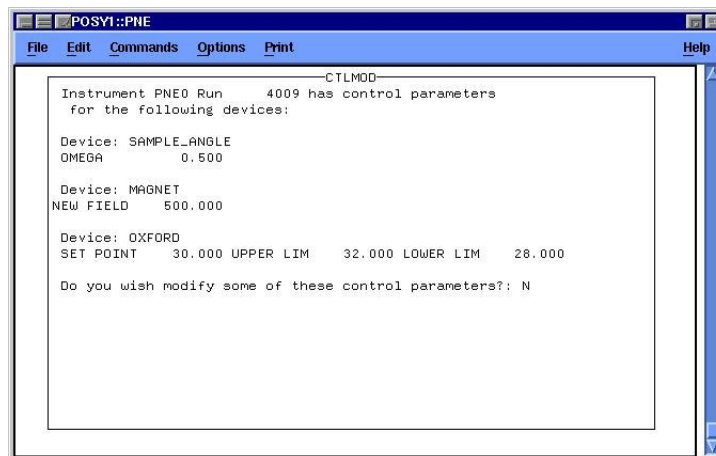
PNE0 RUN      4009: Goyette
FeF2 0.5 Degrees, 500 Oe, 30 Kelvin
PNE0 First started at 10:47:29 on 12-AUG-97 : Total Live Pulses = 0
Total of Monitored Variable in File = 0
PNE0 ACTIVE_OFF last stopped at 10:32:39 on 12-AUG-97.

0 of 720 cycles completed.
0 of 108000 pulses completed for cycle 1.

POSY1_PNE>
```

Modifying Control Parameters

If, after completing the set up a run file, the modification or deletion of a control parameter in that run file is desired, use the command CTL. The program will display the control parameters that have been set for that run file. You have the option of changing the parameters that have been set or removing a control parameter completely. You do not have the option of adding a new control parameter.



```
POSY1_PNE> CTLMOD
Instrument PNE0 Run      4009 has control parameters
for the following devices:

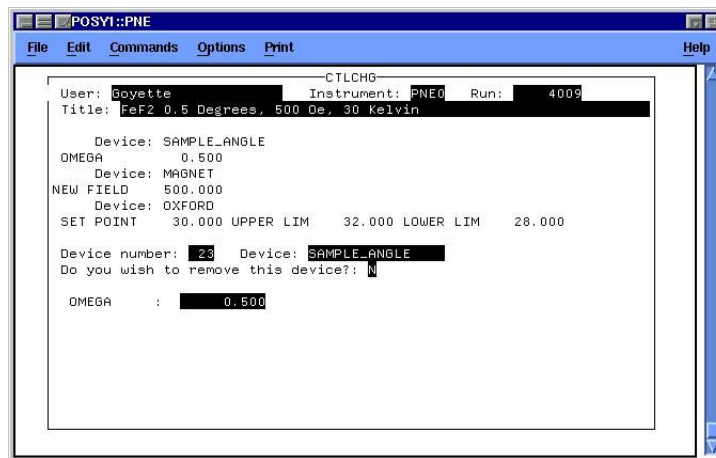
Device: SAMPLE_ANGLE
OMEGA      0.500

Device: MAGNET
NEW FIELD   500.000

Device: OXFORD
SET POINT   30.000 UPPER LIM   32.000 LOWER LIM   28.000

Do you wish modify some of these control parameters?: N
```

The program will go through each control parameter that has been set. It will first ask if you wish to remove the device:



If you answer yes, it will be removed completely. If you answer no, it will ask if you wish to modify the value given for the device. Once the program has completed a device, it will move on to the next device where the process is repeated. When all parameters have been completed, the program will end. If any modifications were made, you should check the new parameters by using PRUN CONTROL.

Reading T0 Pulses Using RDP

You can monitor the number of T0 pulses that have been counted by typing the command RDP. The number of T0 pulses will be displayed:

```
PNE0: NUMBER OF T0 PULSES= 107778
```

Reading Sample Temperature

You can monitor the sample temperature by typing the command RDT. You will receive a menu similar to the menu when using TEMTUR:

```
POSY1_PNE> rdt
*** RDTEMP ***
WHICH TEMPERATURE CONTROL DEVICE DO YOU WANT?
DEVICES AVAILABLE ARE:
  4  DISPLEX
 26  LAKESHORE
 35  OXFORD
 36  TRANSLATOR
 37  LAKESHORE LS340
 38  RECIRCULATOR
ENTER ----> DEVICE NUMBER: 37
PNE0 MEASURING TEMPERATURE WITH LAKESHORE LS340 CON-
TROLLER -- DEVICE 37
PNE0 LAKESHORE LS340 CTL TEMP (1) = 296.58 : MEAS TEMP (2)
= 9998.00
PNE0 LAKESHORE LS340 % OP PWR (1) = 0.00
```



Looking at the Data

This chapter describes methods by which you can look at live data as it is being collected, or at old data stored on the hard disk. Data may be viewed as a function of time of flight (which can be converted to wavelength), or as a function of position channel (which may be converted to angle). This chapter also describes how to obtain hard copies (printouts) of raw data.

Intensity as a Function of Position

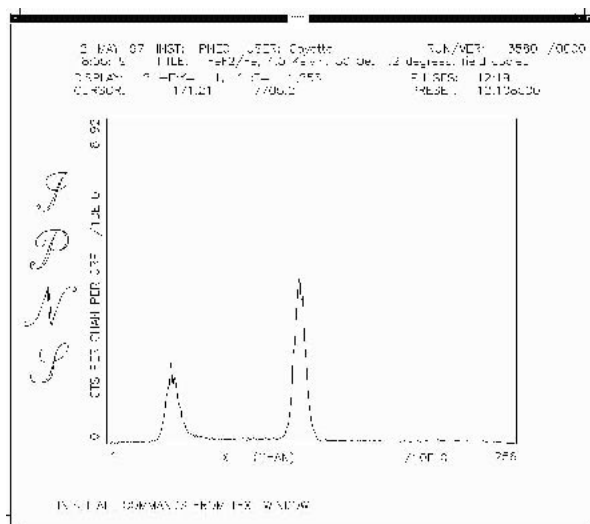
To view live data as a function of position channel for a particular time-of-flight, type:

TYP LH <cr> (stands for Live Histogram)

HST X <cr> (Histogram as a function of position channel X)

TPT time channel # (1-256) <cr> (time channel #corresponding to the time-of-flight of interest, it can be a number from 1 to 256)

DIS <cr> (to display a plot similar to the one below)



When a data display is active, placing the mouse arrow within the data display region will change the mouse arrow to a cursor (+). The cursor position is controlled by the mouse. The cursor is useful in determining channel positions and intensities. Place the center of the + at the position of interest, and read the channel and intensity values from the screen.

To clear screen: Type: CVS <cr>

Intensity as a Function of Time-of-Flight

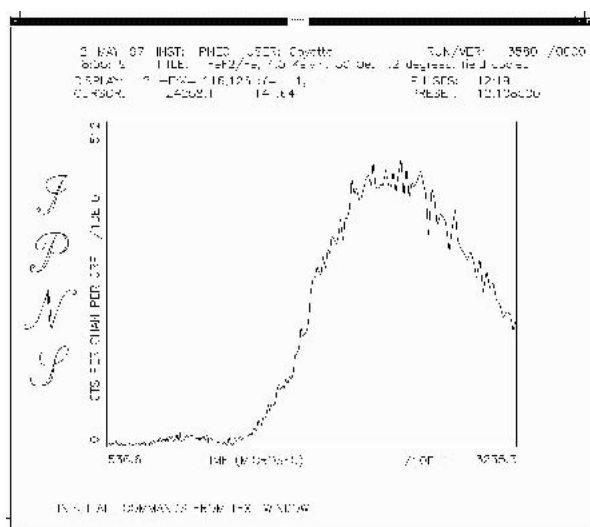
To view live data as a function of time-of-flight for a particular position channel, type:

TYPLH <cr> (if you have typed this already, you don't have to do it again)

HST T <cr> (histogram as a function of time-of-flight)

XPT POSITION CHANNEL # (1-256) <cr> (x position channel of interest, it also can be a number from 1 to 256)

DIS <cr> (to display a plot similar to the one below)



If you want a printed version of any of the plots that appear on the screen, type DPLLOT PS <cr>. The data screen will disappear, and the computer will create a file labeled PGPLOT.PS. Wait about ten seconds and then type:

PRINT/QUEUE=LJ4MV_399 PGPLOT.PS <cr>

and a printed copy of the plot should appear on LJ4MV printer in Bldg. 399. There are several formats in which a plot can be saved (GIF, Post Script, Color Post Script, etc. Consult a member of the POSY1 Staff for more details.)

Rescalling the X and Y Axis of Any Plot

To change the upper and lower limits on either the X or the Y axis use the following procedures:

Changing the Y-Axis

The general commands to change the y-axis are these:

YMIN <cr> to change the minimum of the y-axis.

YMAX <cr> to change the maximum of the y-axis.

YSC <cr> to toggle between automatic and manual scaling.

The computer will indicate which mode you are in after you type YSC. To specify a YMAX and YMIN of your own, you must be in Manual scaling mode. DIS <cr> as always, to display the plot.

EXAMPLE: changing YMAX from the current value to 5000:

YMAX 5000 <cr>

YSC <cr> (to go into manual mode)

DIS <cr>

To return to the automatic scaling, just Type:

YSC <cr> (to go back to automatic scaling mode)

DIS <cr>

Changing the X-Axis

To reset the min and max values for the x-axis, use the following command:

DX NEW XMIN NEW XMAX <cr>

Where NEW XMIN is the desired value of XMIN and NEW XMAX is the desired value for XMAX. EXAMPLE: changing XMIN to 40 and XMAX to 80:

DX 40 80 <cr>

DIS <cr>

To return to the original scaling for the x-axis, use the same command:

DX 1 256 <cr>

Remember that when you plot as a function of position channel you must specify the x-axis in position channels, and when you plot as a function of time-of-flight you must specify the x axis in microseconds. the full time range (also the default range) is from 5,366 μ s to 32,353 μ s. This means that when you switch from x to time of flight you must give the command

DX 5366 32353 <cr>

and when you switch from time of flight to position channel you must give the command

DX 1 256 <cr>.

Looking at Different Spin States

Data for spin up neutrons is stored in Y channel 1, and can be viewed by typing:

```
YPT 1 <cr>
```

as part of the data display commands. Data for spin down neutrons is stored in Y channel 2, and can be viewed by typing:

```
YPT 2 <cr>.
```

Integrating Over Several Channels

You may integrate over several position channels or time channels by specifying the inclusive range of channels when giving either the TPT command or the XPT command.

```
TPT Lower limit Upper limit (when integrating over time channels)
```

or

```
XPT Lower limit Upper limit (when integrating over position channels)
```

For example, lets say you want to integrate from position channel 20 to position channel 105. You would type:

```
XPT 20 105 <cr>
```

When integrating over time channels, you specify the times (in microseconds) over which you wish to integrate.

```
TPT 15000 30000 <cr>
```

and follow all other details from the first example in this chapter

Looking at Old Data

You can also look at data that was collected previously by using the RUNN command to recall a previous run along with the commands to display file data:

```
TYP FD or TYP FH <cr> (File Density, and File Histogram)
```

```
RUNN run# <cr>
```

```
DIS <cr>
```

The commands HST X and HST T may be used as with live runs.

Looking at Monitor Data

Monitor data may be observed in the same way as detector data. The only difference is that you specify TYP L for live monitor data and TYP F for file monitor data.

Reinitializing the Graphics Software

It is sometimes necessary to reinitialize the software that displays a plot on the VAX screen. When this is the case, simply type DINT, which will reinitialize the plotting

package. If this does not work, type DINTALL. After this, you are ready for normal data display procedures.

Operator Messages

Operator messages are sometimes displayed on the screen by the MicroVax. These are usually announced by a change of the screen to half black and half “normal”. Initially, it may appear as if a serious system error has occurred; but this is not the case. Hold the CONTROL key down while pressing the F2 key, located at the upper left of the keyboard will switch back to the regular screen.



Performing a Transmission Run

A transmission run should be performed after any change in the moderator has occurred. In addition, a transmission run should be performed after changes of or modifications to sample holders.

Note: You must decide beforehand whether you will be using a transmission run to analyze data that was collected using the spin analysis mirror. If the transmission run is not to be used on spin analyzed data, the mirror must be out of the beam when the transmission run is collected. If the transmission run is to be used on spin analyzed data, the mirror must be inserted in the beam so that the transmitted beam passes through the silicon substrate of the mirror before reaching the detector.

Setting Up a Transmission Run

To perform a transmission run, move the sample out of the beam. Do this by using the command TRANSLATE, selecting option P, and entering a destination position that is approximately 3 millimeters more positive than the current position. This will translate the sample stage 3 millimeters to the left, (from a viewer's perspective if you are standing at the instrument, facing the cryostat, with your back to the detector). Slide the boron nitride mask to the right until the detector face is unobstructed. Record the current setting of the collimator and then close the collimator to 15 on the dial. Open the beam gate and start a new run using the method discussed later. After the run is completed close the beam gate, move the boron nitride mask to the left until the rightmost edge of the mask is aligned with the rightmost edge of the aluminum track on which the mask slides. Place the sample back in the beam and return the collimation to its original setting. Normal operation is now restored.

Processing a Transmission Run

After the transmission run has finished, it may be analyzed and a TRANS.DAT file may be created by using the POSYIDL2 program. You can only use POSYIDL2 from an X-windows terminal. Different platforms use different programs, which must be configured properly to allow X Window connections. To use the POSYIDL2 program,

log on to one of the IPNS LINUX systems where the POSY1 data is mounted using your LINUX username and password. If necessary, set your display variable to the IP address of the computer where you sitting. Then type POSYIDL2. Click the button labeled POSY1 in the window that appears. You will be presented with the main POSY2IDL window. Select Analyze Data from the Data Analysis menu. Select Spinread from the Analyze Data sub-menu. You will be presented with the Analyze experimental data window:

Run number: 7966

Filename root: trans

Title: 7966,transmission run through cryo+RT inser

With Flipper ☐ Sum Multiple Runs ☐ no ☐ yes

% of delayed neutrons: 0.417 number of files to sum: 1

Pick specular position

Left Bg Start: 0 Left Bg End: 7

Right Bg Start: 65 Right Bg End: 65

Left Signal Start: 27 Left Signal End: 35

Right Signal Start: 1 Right Signal End: 1

Execute Done

Instructions

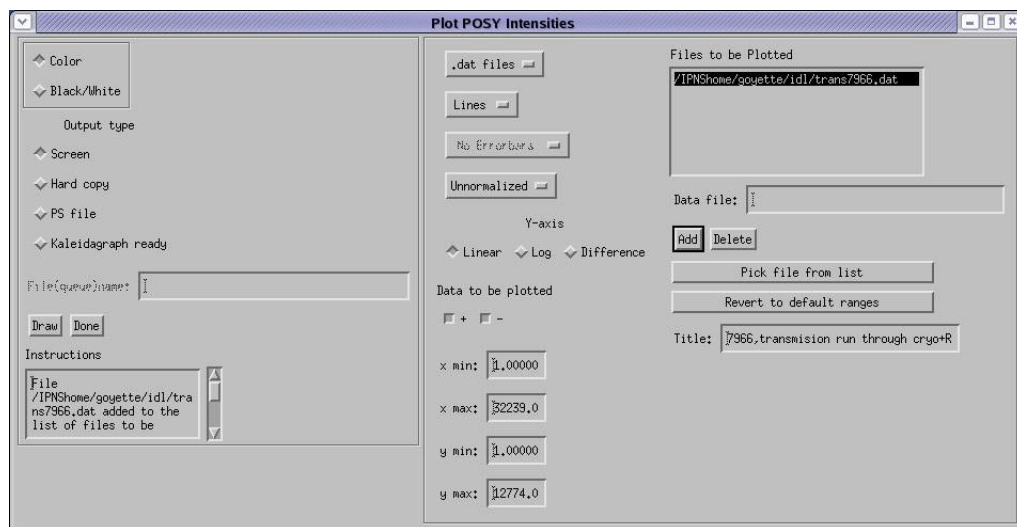
Signal region: green.
Background regions: red. If these values are acceptable press <Execute>. If not,

Type the run number of the run file you wish to analyze into the box labeled Run Number, and press return. You must press return from the Run Number box for the program to correctly read all the information from the run file. Once the run file has been read, the title box in the window will be filled automatically. Also notice that after the program had finished reading the run file, a position histogram of the run file appeared in the main window of POSYIDL. Multiple runs can be summed together, if desired, by selecting the button labeled YES under Sum Multiple Runs. An additional window will appear requesting which run numbers to sum together. The default Run Number box will become inactive when multiple runs are summed together. You can modify the title or any of the other data in the window. In the box labeled filename root type TRANS to signify that this is a transmission file. Click the button immediately below the Title field and select WITH FILPPER. Once the run file has been read in, click the button labeled Pick Specular Position. Now click on the header of the main window (make sure you click on the header and not inside the plot area). Identify the position of the transmitted beam, and click on the center of the peak with the cursor. The program will determine the signal regions (highlighted in green) as well as the left and right background regions (highlighted in red). Move back to the Analyze Experimental Data window and you will notice that the values for signal and background region will now be displayed in the appropriate boxes. If you do not like these values, you may type in your own. In addition, if for some reason (such as you clicked in the plot window when you should have clicked on the header) the program found very bad values for the regions, you can have the program refind the signal and background regions by clicking the Find Specular

Position button again. When you are ready to write a TRANS.DAT file to disk, press the execute button: The instructions box will tell you when the file has been written. The instructions box will also contain messages on how to use the program.

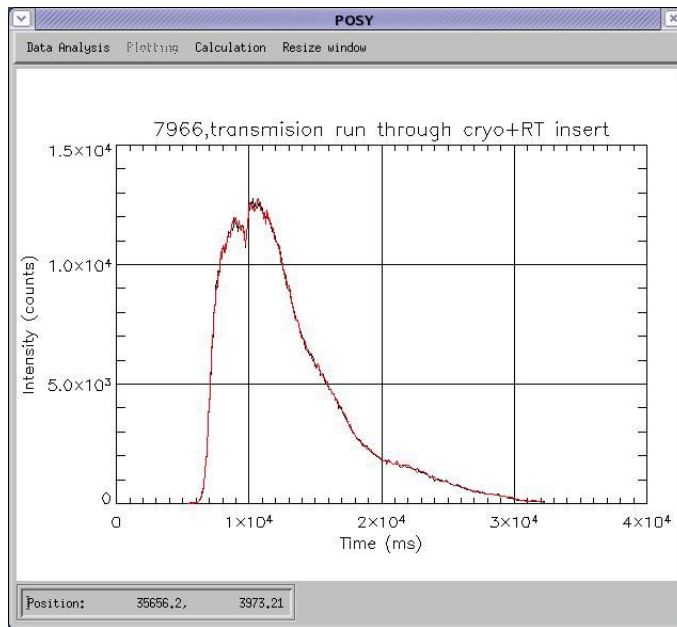
Plotting a Transmission Data File

After generating a transmission data file (a TRANS.DAT file) you can plot the results to the screen or to a printer by using the POSYIDL program and selecting Analyzed Data from the Plotting Menu. You will get the following window:



Click on the topmost button on the left hand side of the window and select Transmission. Type in the name of the file in the box labeled Data File and click the button labeled Add, or use the Pick From List button to see a list of files in the current directory. You can change the directory in which you are looking by typing in the new directory path in the box labeled Path. You can also look at only some files by specifying a filter. Click on the file you want, and press the button labeled OK. The file you clicked on should show up in the Data file box. Click Add to add the file to the Files to be Plotted list. When you are in the Plot POSY Transmission Files window you can specify any combination of points, lines and error bars for the plotting of the data. You can also specify whether you want a color or black and white plot, and whether you want to plot to the screen, to a Post Script file or to a printer (Hard Copy). If you request that the plot be sent to a file, you will have to enter a file name into the box labeled file (queue) name. This box also stores the name of the queue to print to if you request that the plot be sent to a print queue. When you have made all the appropriate selections, press the draw button. The Instructions box will inform you when the plot is finished, and will offer instructions on how to use the program. If you have selected the screen as output, you will get back the main window which contains the plot:

Performing a Transmission Run



When you are done viewing the transmission data file, click the Done box in the Plot POSY Transmission File window. This window will close, but the main window, which still contains the plot will remain. This is normal.



Data Analysis

This Chapter describes how to extract information from the raw data, analyze the raw data and display the results in several different forms. It is generally helpful to have a member of the POSY1 staff on hand at this point until some experience is obtained.

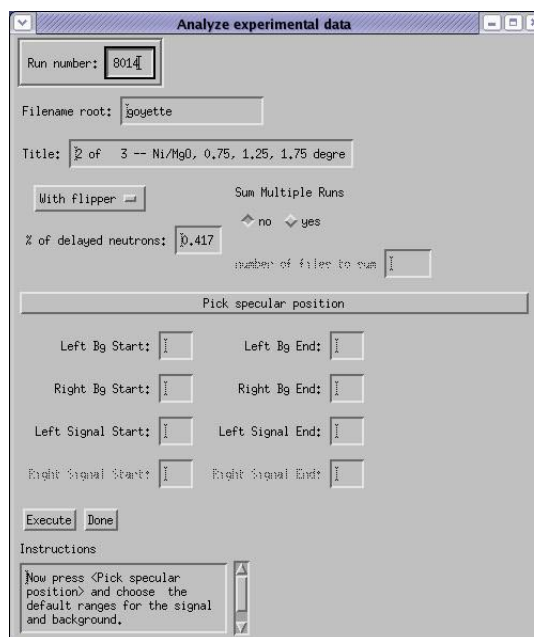
Note: Before generating a reflectivity file, you must identify whether that run was unpolarized (no flipper), with flipper only or with flipper and analyzer. This must be correctly specified in SPINREAD. In addition, the transmission run used to calculate the reflectivity file in SPINREF must be of that same type as the file generated by SPINREAD.

Generating a Reflectivity File

Run files may be analyzed, and reflectivity files generated, using the POSYIDL2 program. POSYIDL2 can only be used from an X-window terminal. Refer to the chapter on Performing a Transmission Run for the procedure on how to bring up a POSYIDL2 window.

Creating a .DAT File Using Spinread

SPINREAD will generate a .DAT data file of raw intensities that have been corrected for background counts and delayed neutron counts. SPINREAD does not calculate the reflectivity. When the POSYIDL main window appears, select Analyze Data from the Data Analysis menu. Select Spinread from the Analyze Data sub-menu. You will get the Analyze Experimental Data window:



Type the run number in the box labeled Run Number, and press return. Remember that you must press return while you are in the Run Number box for the program to work correctly. Multiple runs can be summed together, if desired, by selecting the button labeled YES under Sum Multiple Runs. An additional window will appear requesting which run numbers to sum together. The default Run Number box will become inactive when multiple runs are summed together. Once the program has finished reading the run file, the Title box will contain the title of the run file. Specify the file name prefix (usually the users last name) in the box labeled Filename Root. After the program had finished reading the run file, a position histogram of the run file appeared in the main window of POSYIDL. Click the button immediately below the Title field and select the appropriate run setting. Once the run file has been read in, click the button labeled Pick Specular Position. Now click on the header of the main window (make sure you click on the header and not inside the plot area). Identify the specular position, and click on the center of the specular peak with the cursor. The program will determine the signal regions (highlighted in green) as well as the left and right background regions (highlighted in red). Move back to the Analyze Experimental Data window and you will notice that the values for signal and background region will now be displayed in the appropriate boxes. If you do not like these values, you may type in your own. In addition, if for some reason (such as you clicked in the plot window when you should have clicked on the header) the program found very bad values for the regions, you can have the program refind the signal and background regions by clicking the Find Specular Position button again. When everything is set correctly, press the button labeled Execute to generate a .DAT file. A message will appear in the Instructions box when the reflectivity file is finished. The Instructions box will also contain instructions on how to use the program.

Createing a Reflectivity File Using SPINREF

Once a .DAT file has been generated using SPINREAD, a reflectivity file can be created using a transmission file and the program SPINREF. If a transmission file does not yet

exists, see the instructions on setting up and processing a transmission run. Once a valid transmission file exists, select Analyze Data from the Data Analysis menu. Select Spinref from the Analyze Data sub-menu to get the SPINREF window:

Calculate reflectivities

Output filename: /IPNSHome/goette/id

Reflection filename: /IPNSHome/goette/id1/goette8312.dat

Reflection file information

1 of 2 -- 8312+8313 Ni/MgO (001) annealed. Center at channel: 105.692. File contains 124 bad values. These will be set to zero before normalization. Reflection file with

Transmission filename: /IPNSHome/goette/id1/trans7966.dat

Transmission file information

7966, transmission run through cryo+RT insert. Center at channel: 30.9532. Transmission file with flipper (2 sets of data).

Detector Translation (in inches): 12.0000

Transmission Channel: 7.5000

Number of Channels per Degree: 106.667

Reflection Angle: 0.742406

Title: 1 of 2 -- 8312+8313 Ni/MgO (001) annealed

Instructions

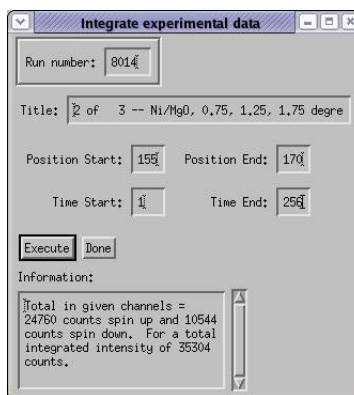
REF+ is black, REF- is red. If all of the parameters look correct press the "Write output" button to write the data to a file. The default filename is the reflection filename

Enter the name of the reflection filename (the .DAT file). Enter the transmission filename. The filenames for reflection file and transmission file can be typed in directly, or the PICK buttons for each may be used to see a list of files from which to choose. You can change the directory you are looking in by typing in the new directory path in the box labeled Path. You can also look at only some files by specifying a filter. Enter the output filename. This name will usually be the reflection filename with the suffix .OUT instead of .DAT. If no output filename is entered, the default name of spinref.out will be used. Click the READ FILES button. The files will be read, and some information from those files will be displayed in the information boxes. The sample angle will be calculated and displayed in the box labeled Reflection Angle. SPINREF will use this angle to generate the reflectivity -vs- k_z data file that will become the .OUT file. This angle can be modified after the initial angle calculation is made. The new angle will be used to calculate k_z . If a value of 0.0 is entered for the angle, the .OUT file will be written as reflectivity -vs- wavelength instead of reflectivity -vs- k_z . Other settings, such as scale factors and hurdle factors should be discussed with a member of the POSY1 staff. Once the files have been read in, The reflectivity -vs- wavelength plots will appear in the POSYIDL main window. SPINREF will not display the k_z reflectivity. When you are ready to save the .OUT file, click the button labeled WRITE OUTPUT. SPINREF waits until the WRITE OUTPUT button is pressed before calculating k_z values, if necessary. Click the DONE button when finished.

Calculating Polarization using POL

The Average polarization of a run file can be calculated using the program POL. Within

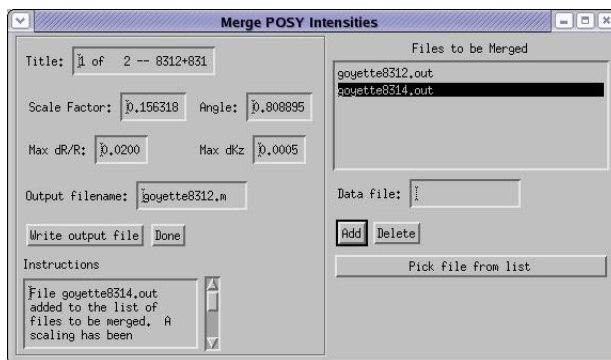
a specified time-of-flight range, POL will add together all the spin up counts and divide the total by the sum of all the spin down counts. The result is defined to be the average polarization for that time-of-flight range. To calculate the average polarization for a run file, select Analyze Data from the Data Analysis menu and select POL from the Analyze Data sub-menu:



Enter the data file name (it must be a .DAT file) and press return, or use the PICK button. The polarization will be calculated for the entire range of times. A plot of the data will appear in the POSYIDL Main Window. To select a different range of times, enter new start and end times. The calculated polarization and the plot will automatically update. Click Done when finished.

Merging Reflectivity Files

You can merge up to four reflectivity files using the POSYIDL program. Select Merge Reflectivity Files from the Data Analysis menu. You will get the Merge Data Sets window:



Type in the name of the files to be merged in the box labeled Data File, or use the Pick From List button. Enter the data files one at a time and then press the button labeled ADD to add them to the list. Enter the files in order from lowest angle data file to highest angle data file. These data files must be .OUT files. As each file is entered, a scaling factor and angle will be automatically calculated for each file and displayed in the appropriate box. You can view each scaling factor and angle by clicking on individual file names in the list. scaling factors and angles may be modified simply by typing in a new value in the box labeled Scaling Factor. As each file is entered into

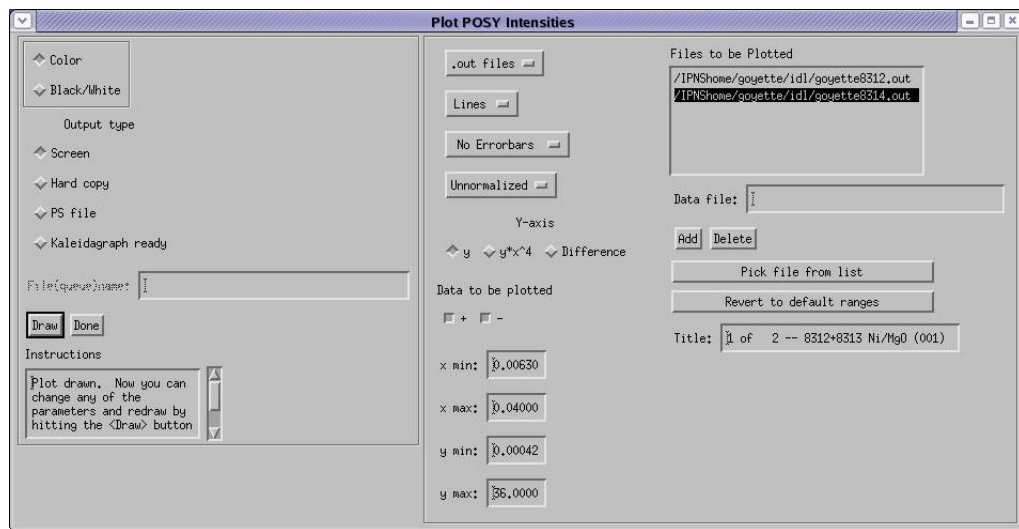
the list, it will be merged with the previous file. Modifications to any scaling factor and angle can be observed immediately in the plot window. Changes in dR/R and dKz will affect how densely populated the points are in the merged file. When a good merged is obtained, enter an output file name and click the button labeled Write Output File. The name of the merged file will be automatically generated from the name of the normalization file (goyette8300.out, for example) with the “.out” extension changed to “.m” (goyette8300.m) In addition to the merged file, output files compatible with the fitting package Parratt32 will be generated. Separate files will be written for spin up and spin down data. Parratt compatible filenames will be generated in a similar manner to the full merged filename, except that the name will include a “p” or an “m” to distinguish plus (up) from minus (down) spin states, and a “.dat” extension, which is required for Parratt32 (so goyette8300p.dat and goyette8300m.dat in our example).

Plotting Experimental Data

You can plot (to the screen, to a file or to a printer) reflectivity files (.DAT, .OUT and .M files) and run files (.RUN files) using the POSYIDL program.

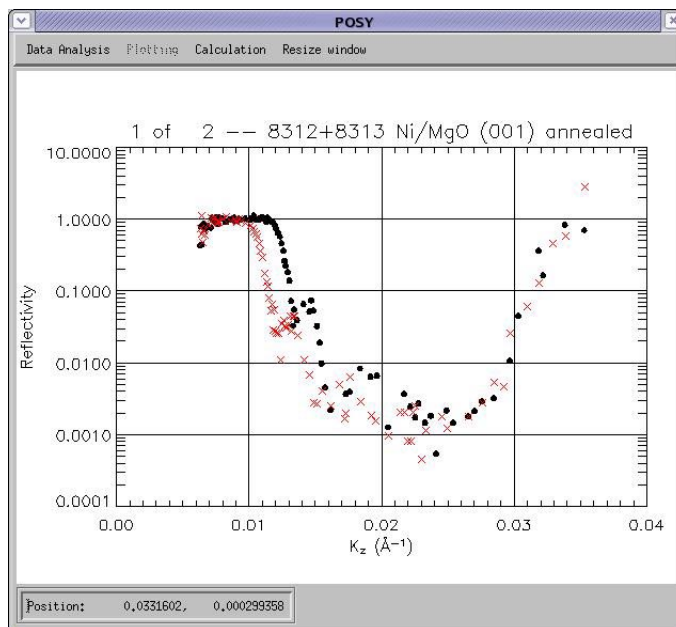
Plotting Reflectivity Files

To plot a reflectivity file, select Analyzed Data from the Plotting Menu. You will get the Plot POSY Reflectivity File window:



Click on the topmost button on the left hand side of the window and select Reflectivity. As we did when we were plotting a transmission run, type in the name of the file in the box labeled Data File, or use the Pick From List button to see a list of files in the current directory. When you are in the Plot POSY Reflectivity Files window you can specify any combination of points, lines and error bars for the plotting of the data. You can also specify whether you want a color or black and white plot, and whether you want to plot to the screen, to a Post Script file, to a file that can be read easily by KaleidaGraph or to a printer. You can also specify whether you want to plot R , or R_k^4 . You can also specify which polarization cross sections you wish to plot ($++$, $--$, $+-$, $-+$). The $+-$ and $-+$ cross sections will be de-sensitized if the spin analysis mirror was not present for that

run. When you have made all the appropriate selections, press the Draw button. The Instructions box will inform you when the plot is finished, and will offer instructions on how to use the program. If you have selected the screen as output, you will get back the main window which contains the plot:

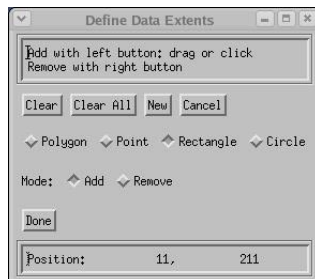


When you are done viewing the transmission data file, click the Done box in the Plot POSY Reflectivity File window. This window will close, but the main window, which still contains the plot will remain. This is normal.

Plotting Contours of Run Files

You can generate contour plots of run files by using the POSYIDL program. Select contour from the Plotting menu. You will get the Contour Plot POSY Data window:

As always, type the run number in the box labeled Run Number and press return, making sure you press return while you are in the Run Number box. When the program has finished reading the run file, a plot of the data will appear in the main POSYIDL window. This plot is unlabeled and displayed only in channel numbers. To display the plot in theta and lamda, press the button labeled Define Data Extents. The Define Data Extents window will appear:



You will need this window in a moment. First, switch back to the main window which contains the unlabeled plot. Using the cursor click and drag a box around the data you wish to plot. Click DONE on the DEFINE DATA EXTENTS window once the box is correct. If the box is not right, redraw it. After you click Done, the define data extents window will come back again. Switch back to the plot window. Draw a box around the specular reflection just like you did for the data. Click done on the Define Data Extents window again. Then switch back to the Contour Plot POSY Data window and click the button labeled Draw. The plot in the main window will be expanded so that the range you selected with the cursor box will fill the plot window, and the axes will be labeled in theta and lamda. You may modify the contour to display in Q_x and Q_z instead of theta and lamda. You may also generate a color contour, where the contour levels are lines or filled. You may direct the output to the screen, to a printer or to a Post Script file. The contour may be normalized to a transmission run. The number of contour levels can be changed and you can always type in the data extents by hand if you would prefer to do so. Contour plots may be normalized to a transmission run by clicking the Normalized button and identifying the appropriate transmission file. Detector translation should not be modified without first consulting a member of the POSY1 staff. When you are finished, press the button labeled Done on the Contour Plot POSY Data window. The window will disappear, and you will be back to the main POSYIDL window.



Data Fitting and Modeling

This chapter describes how to create and modify data model files, generate calculated data sets, and compare them to experimental data sets using chi squared fitting. It is necessary for you to have some experience using the program LAYERS before using these fitting routines on your own. If you do not have the necessary experience, obtain the assistance of a member of the POSY2 staff to fit your data.

Fitting Data Using LAYERS

The fitting routines LAYERS or CALPAK can be run from the POSYIDL program by selecting Layers from the Calculation pulldown menu. You will get the RUN Layers window:

Scattering Density R(b/v) (10 ⁻¹² cm/A ² 3)	Magnetic Im(b/v) (Gauss)	Magnetic Parallel (Gauss)	Magnetic Normal (Gauss)	Layer Thickness (Å)	Roughness Halfwidth (Å)	Tag
0.0652500	0.0000000	0.0000000	0.0000000	150.000	15.000	0.000
0.0207300	0.0000000	-0.0412400	0.0000000	80.000	15.000	0.000
0.0257000	0.0000000	0.0000000	0.0000000	75.000	10.000	0.000
0.0914600	0.0000000	0.0082400	-0.0206500	120.000	5.000	0.000
0.0652500	0.0000000	0.0000000	0.0000000	150.000	5.000	0.000
0.0207000	0.0000000	0.0000000	0.0000000	9999.000	0.000	0.000

Instructions:
Edit any of the Parameters and hit execute.

The program will automatically read the current PM.DAT file and display it in the text window. The values of Kmin, Kmax, deltaK and Kstep are displayed separate from the rest of the data in PM.DAT, but will still be written the old way when PM.DAT is updated. Type in the experimental file name in the box labeled Experimental Filename.

Select the appropriate button to choose either LAYERS or CALPAK. When choosing CAPLPAK, also specify the desired cross section. Make any desired modifications to the profile and press the button labeled Calculate. The PM.DAT file will automatically be saved, and the current Fortran version of LAYERS will be executed. The experimental data file and the calculated file will be displayed in the main POSYIDL window. As before, the experimental file will be plotted using the + symbol, while the calculated file will be plotted using a line.



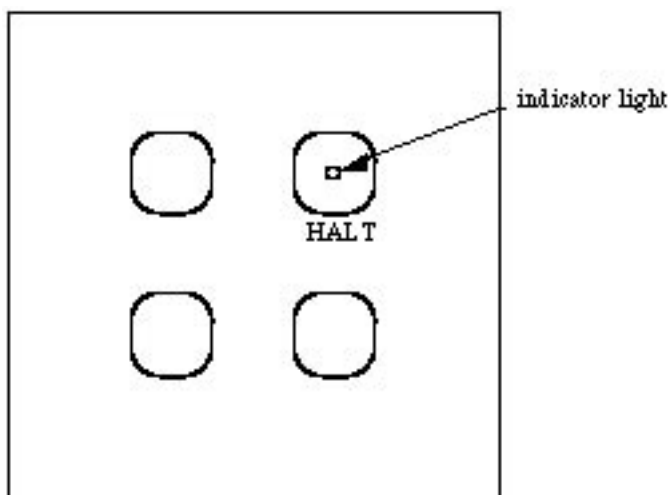
Rebooting the MicroVax

NOTE: NEVER PERFORM ANY TYPE OF REBOOT WITHOUT FIRST OBTAINING PERMISSION FROM A MEMBER OF THE POSY1 STAFF.

Performing a Full System Reboot

If the microvax is not on, or it develops a problem and must be rebooted, here is the procedure:

- 1 Press the Halt button on the front of the computer (see figure one). The red indicator light should come on.



- 2 Press the Halt button again. A >>> should appear on the screen.
- 3 type B DUA0 <cr>. (<cr> means carriage return). The last character is a zero.

When the booting procedure has been completed successfully The logon prompt will be displayed.

Performing a Minimal System Reboot

In the event of an error which occurs due to a lack of disk space, it may be necessary to perform a minimum boot of the system in order to free up enough disk space for the

system to run properly. If there is insufficient disk space, the system will not be able to run properly. It will disable logins, and will not perform a normal boot correctly. When this happens, follow the following instructions for a minimal system boot, login to the user account and remove enough files so there is at least 20000 blocks of free disk space, and then perform a normal boot.

1. Press the Halt button twice to get to the >>> prompt.
2. Type B/1 DUA0 <cr>.

After a few moments, you will receive the SYSBOOT> prompt.

3. Type SET STARTUP_P1 "MIN" <cr>.

You will get the SYSBOOT> prompt again.

4. Type CONTINUE <cr>.

The computer will now perform a minimal system boot. Under the minimal system, no windows will be created. When the boot is completed, the window will remain just as it was during boot. Press the Return key a few times to get the Username: prompt and log onto the computer giving the username and password for the user account. Once the login is complete, purge the entire user account by issuing the command PURGE [PNE...] <cr>. Once the purge is completed check the free disk space by typing SHOW DEVICE DUA0: <cr>. You will receive the response:

Device	Device	Error	Volume	Free	Trans	Mnt
Name	Status	Count	Label	Blocks	Count	Cnt
POSY2\$DUA0:	Mounted	0	VAXVMSV055	38493	219	1

If the number of Free Blocks is less than 20000, it will be necessary to delete some files until there are 20000 free blocks. Run files are backed up regularly. In addition, a complete system backup is performed once a week, and incremental backups are performed every day. As a result, anything that is removed should be available for restoration from tape. Delete the oldest run files (those with the lowest run number) available, since these are the largest files and are sure to have been backed up. Make a note of the files that were deleted.

Once there is enough space on the disk:

1. Press the Halt button twice to get to the >>> prompt.
2. Type B/1 DUA0 <cr>.

you will get the SYSBOOT> prompt.

3. Type SET STARTUP_P1 "" <cr>. There is nothing between the quotes.

you will again get the SYSBOOT> prompt.

4. Type CONTINUE <cr>.

The computer will now perform a full system boot.



Analyzer Instructions

Instructions on using the POSY1 analyzer

Using the analyzer.

To use the analyzer, it will need to be translated into the beam and rotated to the correct position. The analyzer is a beam splitter, one of the spin states goes straight through, and the other one is reflected from the analyzer. Unfortunately the analyzer is not curved nor is it long enough, so the beam that transmits through the analyzer is often contaminated with beam that misses the analyzer completely. That is why we concentrate on the beam being reflected from the analyzer. For the optimal orientation of the analyzer with respect to the beam that is to be analyzed, there should be an angle θ_0 equal to 0.6 degrees between the two. To determine this angle use:

$$\theta_0 \text{ (deg)} = (X_R - X_T) / 38.6.$$

X_R is the position channel for the beam being reflected by the analyzer, and X_T is that of the beam going through the analyzer, which is also that of the beam being analyzed. It follows that there should be a difference of 23 channels on the detector in order to get an angle equal to 0.6 degrees.

To move the analyzer in the beam:

Type “analyzer”

Select “p”

Type “in”

Type “0” for the mirror angle θ . This angle θ refers to that of the sample with the incident beam. When analyzing the direct beam, the sample angle $\theta = 0$.

The analyzer will then automatically rotate a total angle equal to:

$$\varphi := 2\theta + \theta_0.$$

As the default it will use a value of 0.6 deg for θ_0 , which is the desired value.

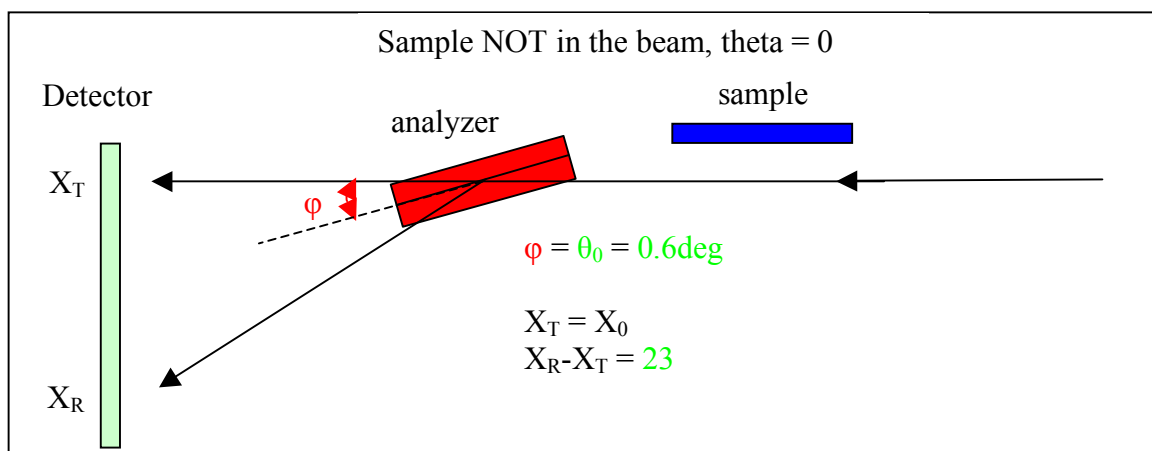
Additionally the analyzer will translate to a value that is calculated based on the provided value theta. Normally this will put the analyzer in the right position, but it can be wise to do some fine tuning. When $\theta = 0$ (analyzing the direct beam, $T_A = 0\text{mm}$), when the analyzer is moved “out”, $T_A = -14\text{mm}$.

To move the analyzer out of the beam:

Type “analyzer”

Select “p”

Type “out”



Fine tuning of the analyzer position:

If you see a reflection you should first adjust the rotation of the analyzer. Using the equation above you can determine how far off from 0.6 degrees you are and then determine how much to change ϕ . Then:

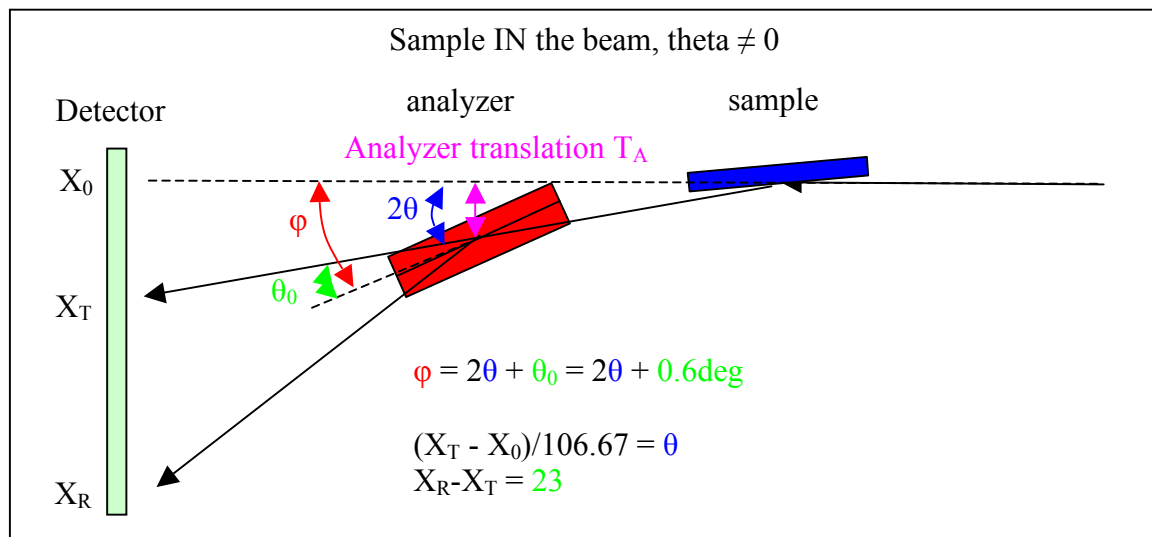
Type “analyzer”
 Select “d” (for diagnostics)
 Select “r” (for rotate)
 Type in the new value of ϕ

Now that the analyzer has the right angle, you will want to maximize the intensity of the reflection (at channel X_R) by translating the analyzer. Do this while looking at the spin up (ypt 1). To translate:

Type “analyzer”
 Select “d” (for diagnostics)
 Select “t” (for translate)
 Type in the new value for the translation T_A .

For each value of T_A record the intensity of the beam reflected by the analyzer by integrating over a fixed number of channels. The position X_R will also change with T_A , but that doesn't matter as long as you keep determining the intensity over a fixed number of channels there the peak is. In other words the number of channels you integrate over stays fixed, but the exact channels change.

When you have found the value of T_A with the highest intensity, translate back to this position. Now check again the value of the analyzer angle θ_0 , if it is 23 channels (0.6deg). If not, again change the angle ϕ as above until this is the case.



Recalibrating the default the analyzer positions (INSTRUMENT PERSONEL ONLY):

After the fine tuning is done for one or more values of the sample angle, a systematic offset for the optimal values of ϕ or T_A might be found, compared to the default settings calculated based on a given value of the sample angle θ .

To change the default values for the rotation (i.e. recalibrate the absolute angle of the mirror), leave the analyzer in the position as optimized for a certain angle θ . Then:

Type “analyzer”
 Select “c” (for calibrate current mirror angle)
 Type in the true sample angle θ

To change the default value for the translation, translate to the optimal position for the analyzer for analyzing the direct beam (without the sample). On the caliper close to the analyzer, press on “zero”, in order to redefine this zero value.

OR, if optimized with a sample at various angles plot T_A versus $\tan(2\theta)$, which should be a straight line. Solve for the parameters D and T_{A0} in

$$T_A = D * \tan(2\theta) + T_{A0}$$

D should be around 790mm, Ideally $T_{A0} = 0$.

Translate the mirror to T_{A0} and press “zero” button on the caliper.

Type “analyzer”
 Select “o” (for changing other parameters)
 Keep pressing return until the line starting with : STOA DIST ?
 Type in the value of parameter D
 Keep pressing return until you get the menu again.



Neutron Reflection

The material in this chapter was taken from *Neutron Reflection Studies of Polymer-polymer Interfaces*, DG Bucknall and JS Higgins, in *Polymers and Surfaces. A Versatile Combination* ed H Hommel, Research Signpost 1998.

Introduction to Reflection Theory

Polymer interfaces and surfaces are an area of study that has always been important since they impinge on all aspects of polymer science. The interfacial behaviour is fundamental to the bulk properties of the system, and consequently the applications of the polymers. For instance adhesion, miscibility, compatibilisation, welding, surface wetting, blend morphology, mechanical strength, etc, are all controlled by interfacial behaviour. The characterisation and understanding of these properties on a microscopic level are therefore of vital importance.

Numerous techniques have been applied to investigate the surface and interfacial behaviour of polymers. However, the accuracy demanded of the experimental data in order to obtain useful interfacial information requires specialist techniques and the associated equipment that goes with them. One technique which has seen an explosion in popularity within the polymer community over the last ten years is specular neutron reflectivity. The reasons for this popularity are partly to do with the development and advances in the reflectometers, but are mostly associated with the simple fact that neutron reflectivity is often the only technique that can answer the questions that are being posed by the polymer interface community. The wealth of information that can be obtained by a well designed experiment would have been difficult to comprehend ten years ago. The object of this contribution is to demonstrate the power of the technique and why it is worth going to one of the handful of neutron sources around the world where neutron reflectometers are operated.

Neutron reflection provides the composition variation normal to the surface of the polymer film, with an accuracy on a sub-nanometer length scale. This kind of information is also given by X-ray reflectivity and to some degree also by light ellipsometry, however the neutron scores over these other radiation sources in two important respects. Firstly, there is the favourable difference in scattering properties between elements and isotopes of elements. With X-rays contrast between different regions of the sample is provided by the electron density difference between molecules, whilst for light it is the refractive index. The analogue for neutrons is the neutron scattering cross section. Across the periodic table, this scattering cross section appears

to be not only of random magnitude, but also of sign and varies even between isotopes of a specific element. For the polymer scientist the neutron is of particular interest because hydrogen and deuterium have vastly different neutron cross sections. This means that with a little careful chemistry deuteration can open up a huge area of unique experimental possibilities. Deuteration is therefore an ideal labelling technique to highlight whole molecules or specific sections of the polymer and produce a large contrast variation with the normal hydrogenous molecules around it.

The second major advantage is that the neutron can penetrate through many engineering materials. This allows the use of neutrons in complex sample environments if the situation warrants it, without the worry of absorption, which is a major problem for light and X-ray techniques. The energies of the thermal neutrons used for reflectivity, make the technique non-destructive, allowing repetitive measurements on one sample to be made, for instance to look at changes after successive annealing processes. The neutrons do not require high vacuum containment which allows samples to be measured under normal atmospheric conditions. Because of these advantages there should be little wonder at the huge growth in the application of neutron reflectivity to the investigation of polymer interfaces and surfaces. However, it must always be remembered that although neutron reflectivity is an extremely powerful technique, it should be used in conjunction with other techniques to fully exploit its potential.

Reflection Theory

It is the wave behaviour of neutrons that has allowed a close analogy with classical optics. With only a few minor modifications, neutron waves are able to reflect, refract and interfere following all the standard optical rules [1]. A full discussion of the theory of reflection is not appropriate at this point since this has been covered extensively in the literature [1,2]. However, it is necessary to include a brief description of the essential points since an understanding of them is vital in the interpretation of reflectivity data.

At this point it is necessary to define a few concepts. An interface is defined as the position between media of 2 different refractive indices. The neutron refractive index of medium j is given as:

$$n_j = 1 - \frac{\lambda^2 N_d b}{2\pi} + \frac{i\lambda N_d \sigma_a}{4\pi} \quad (1)$$

where N_d is the atomic number density, b is the coherent scattering length, the product $N_d b = \rho_z$ is the scattering length density, σ_a is the adsorption cross-section and λ is the neutron wavelength. Strictly then the refractive index is a complex number. For most materials, however, with the exception of those containing the elements, Li, B, Cd, Sm, or Gd, the adsorption cross-section is effectively zero, i.e. $\sigma_a \gg 0$, and therefore for most polymers Equation 1 reduces to:

$$n_j = 1 - \frac{\lambda^2 N_d b}{2\pi} = \frac{\lambda^2 \rho_z}{2\pi} \quad (2)$$

The neutron scattering length b is a nuclear property describing the interaction between the neutron and the atomic nucleus [3]. The value of b varies randomly across the periodic table and also between isotopes of the same element [4]. Here neutrons present a big advantage over other radiation such as light or X-rays, with isotopic substitution providing enhanced contrast between two otherwise equivalent species. The large difference in b between ^1H and ^2H (D) is of particular usefulness to the polymer scientist, as will be discussed in a little more detail later.

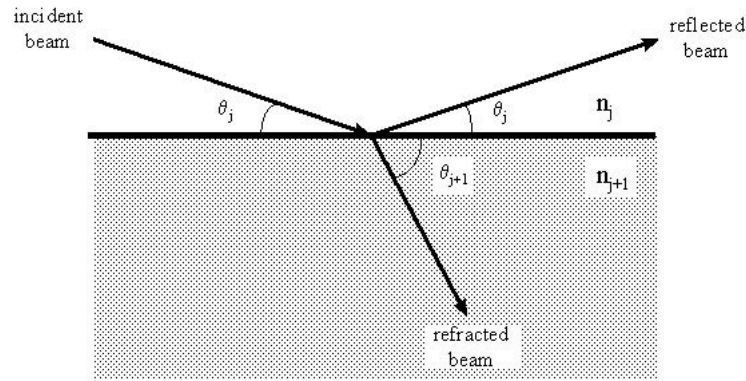


Figure 1: Schematic diagram of an incident neutron beam on a flat interface between two media, showing the reflected and refracted beams.

Figure 1 shows a neutron beam incident on an interface between two infinitely thick media with refractive indices n_j and n_{j+1} , where the interface is also infinitely sharp. The beam can be either reflected, at an angle, θ_j , equal to that of the incident beam, or refracted with an angle according to Snell's Law [5]:

$$n_j \cos \theta_j = n_{j+1} \cos \theta_{j+1} \quad (3)$$

Taking the case where the interface is between a vacuum, or essentially air, ($j=0$) and a liquid or solid ($j+1=1$), then n_0 is 1 and Equation 3 reduces to

$$\cos \theta_1 = n_1^{-1} \cos \theta_0$$

At a critical angle, $\theta_0 = \theta_c$, the angle of refraction becomes zero ($\theta_1 = 0$) and thus $\cos \theta_c = n_1$. For values of $\theta_0 < \theta_c$ total external reflection of the incident beam is observed with only an evanescent wave refracting into the sample. For values of $\theta_0 > \theta_c$ external reflection of the incident beam is accompanied by refraction into the sample.

The reflectivity of an incident beam on an infinitely sharp interface as described above is defined as the ratio of the reflected to incident beam energies [5]. A similar expression for the ratio of the refracted to incident beam energies defines the transmissivity, T . The reflectivity has both parallel and perpendicular components with respect to the interface. It is only the perpendicular reflectivity component that is of importance to the specular reflection of neutrons, since it contains the information about the concentration-depth profile.

The perpendicular reflectivity, R , is determined using the Fresnel reflection coefficient, $r_{j,j+1}$, for the interface described in Figure 1, which is defined:

$$r_{j,j+1} = \frac{n_j \sin \theta - n_{j+1} \sin \theta_{j+1}}{n_j \sin \theta + n_{j+1} \sin \theta_{j+1}} \quad (4)$$

The reflectivity is then simply given by:

$$R = r_{j,j+1} r_{j,j+1}^* \quad (5)$$

where the asterisk (*) denotes the complex conjugate. Since there is no complex component to the reflectivity from most materials then this simplifies to:

$$R = r_{j,j+1}^2$$

The reflectivity from this sharp interface is characterised by the critical angle, θ_c , indicated by the point at which the reflectivity drops from unity, and is often referred to as the critical edge. The wavelength at which this occurs is given by:

$$\theta_c \approx \lambda \left(\frac{Nb}{\pi} \right)^{0.5}$$

Since the neutron momentum transfer is related to the incident angle and wavelength by the relationship

$$q = 2k = \frac{4\pi}{\lambda} \sin \theta \quad (6)$$

it is also possible to derive $r_{j,j+1}$ and hence R in terms of q . Therefore, the reflection coefficients are given by:

$$r_{j,j+1} = \left(\frac{q_{z,j} - q_{z,j+1}}{q_{z,j} + q_{z,j+1}} \right) \quad (7)$$

where $q_{z,i}$ are the component of the momentum transfer perpendicular (normal) to the surface. Since:

$$q_{z,i} = 2 \left(\frac{q_z^2}{4} - \frac{q_c^2}{4} \right)^{0.5}$$

where q_z is defined by Equation 6, and q_c is the critical value of $q_{z,i}$ below which total reflection occurs. Substituting Equation 7 into Equation 5 gives the reflectivity for a sharp interface between two bulk media as:

$$R(q) = \left[\frac{1 - \left[1 - \left(\frac{q_c}{q_z} \right)^2 \right]^{1/2}}{1 + \left[1 - \left(\frac{q_c}{q_z} \right)^2 \right]^{1/2}} \right]^2 \quad (8)$$

This is also known as the Fresnel reflectivity and is often denoted as R_f . A calculated Fresnel reflectivity is shown in Figure 2(a). The region of total reflection below q_c is clearly visible. The drop off in reflectivity at $q_z > q_c$ can be shown to be proportional to $1/q_z^4$, at large values of q_z [2, 6] which is identical to the behaviour observed in systems which have infinitely sharp interfaces and scattering that obeys Porod's Law.

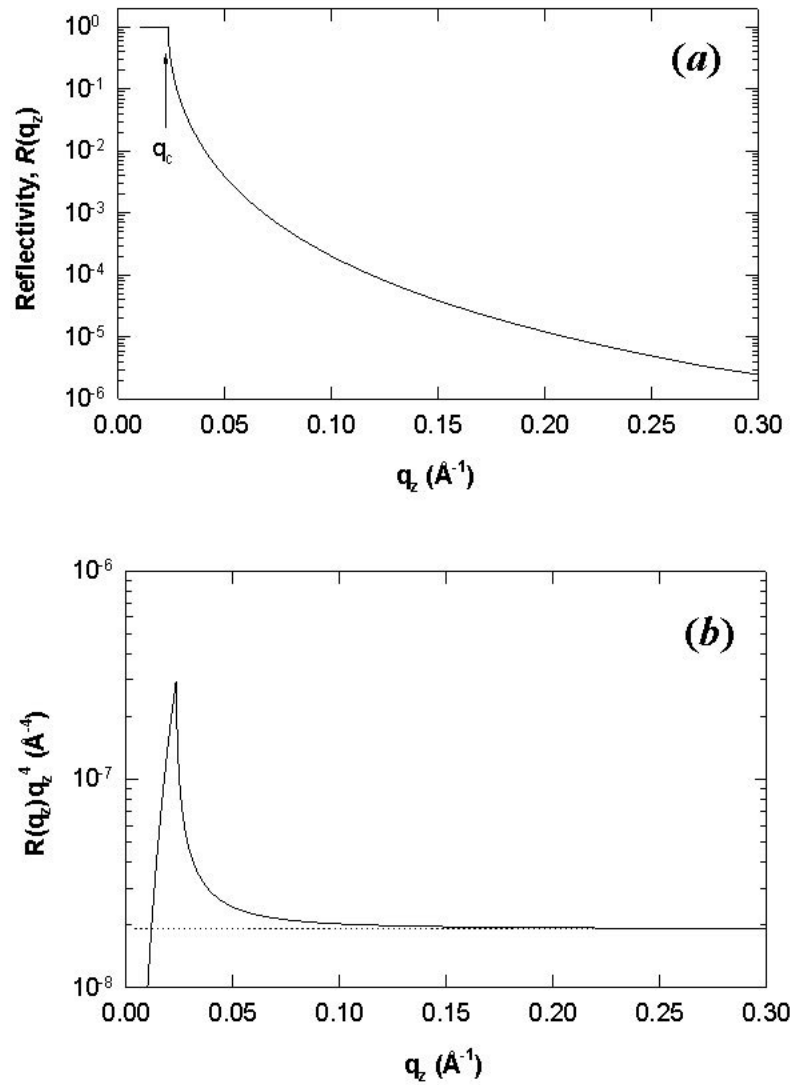


Figure 2: (a) Calculated Fresnel neutron reflectivity profile from an infinitely sharp interface between two media obtained from Equations 4 and 5 or 8. Total reflection ($R=1$) occurs for $q_z \leq q_c$. (b) Reflectivity multiplied by q_z^4 , showing the asymptotic limit reached at large values of q_z .

One interesting result is that the plot of $R(q_z)q_z^4$ asymptotes to a limiting value which is proportional to the square of the reflectance at the air-media interface.

With these equations in mind, the problem of a sample containing m discrete layers can now be addressed. The situation is shown in Figure 3, where the $(m+1)$ layer is the substrate and 0 is defined as the medium surrounding the sample, i.e. air. The reflection coefficient for the sample is calculated by firstly considering the coefficient between the substrate and the bottom layer, $r_{m,m+1}$, i.e. between the $(m+1)^{\text{th}}$ and m^{th} layers, which is simply given by Equation 4 where $j=m$. The reflectivity coefficient between the $(m-1)^{\text{th}}$ and m^{th} is then given by:

$$r'_{m-1,m} = \frac{r_{m-1,m} - r_{m,m+1} \exp(2i\beta_m)}{1 + r_{m-1,m} r_{m,m+1} \exp(2i\beta_m)} \quad (9)$$

where $r_{m-1,m}$ is again given by Equation 4, and the prime (') indicates that internal reflections have been taken into account, in this case between the $m-1,m$ and $m,m+1$ interfaces. A phase factor, β_m , has also been introduced and represents an optical path length term for the m^{th} layer, such that:

$$\beta_m = \left(2\pi/\lambda\right) n_m d_m \sin \theta$$

where n_m and d_m are the refractive index and thickness respectively of layer m , and θ is the angle of incidence. The reflectance, $r'_{m-2,m-1}$, is calculated in a similar way, and the process continued in a recursive method for each successive layer until the reflectivity coefficient for the top layer, $r'_{0,1}$, is obtained. The reflectivity, R , is then given by substituting $r'_{0,1}$ into Equation 5.

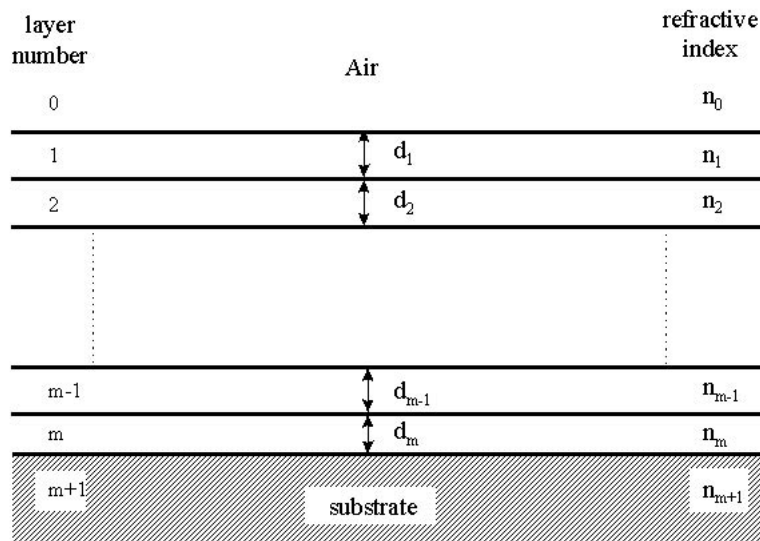


Figure 3: Schematic diagram of a sample composed of m discrete layers each with a thickness, d_i , and refractive index, n_i .

For a double layer sample Equation 9 can be applied together with Equation 5, to give the reflectivity profile shown in Figure 4(a) for a deuterated polystyrene (dPS) layer on a poly(methyl methacrylate) (PMMA) layer on a silicon substrate. The minima and maxima observed are characteristic of the layer thicknesses. When the neutron wavelength satisfies the Bragg condition $\lambda = 2d \sin \theta$, constructive interference occurs. This leads to $d_i = 2\pi/\Delta q_i$, where Δq_i is the difference between successive minima. Therefore the high frequency interference fringes are from the thicker layer, in this case 600 nm, and the low frequency fringes from the thinner layer of 200 nm thick. The reflectivity in the R_q versus q plot as in Figure 4(b) is seen to asymptote to a limiting value at high q values, which is proportional to the sum of the squares of the reflectances at each of the interfaces, given by:

$$Rq_z^4 = 16\pi^2 \sum_{i,j} (\rho_i^2 - \rho_j^2) \quad (10)$$

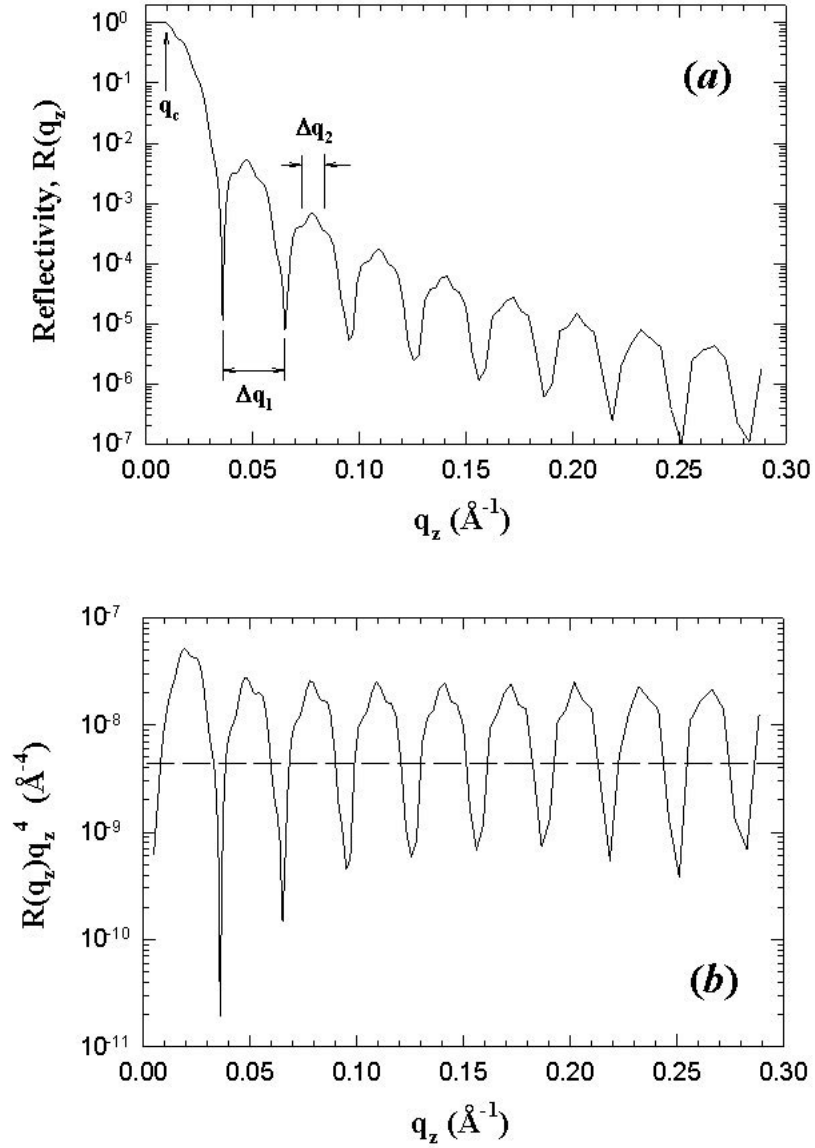


Figure 4: (a) Calculated reflectivity profile of a bilayer of dPS on PMMA on Si using Equation 5 and 9. (b) Reflectivity plotted as R_q^4 versus q , showing that the reflectivity oscillates about an asymptotic value (dashed line).

This approach to calculating reflectivity is exact, but extending it to multilayer systems proves to be extremely cumbersome mathematically. For this reason many people have favoured a more general solution and perhaps the most versatile and widely used among these is that of Abeles [7]. This method defines a characteristic matrix for each discrete layer so that for the m^{th} layer the corresponding matrix is:

$$C_m = \begin{bmatrix} \cos \beta_m & -(i/\kappa_m) \sin \beta_m \\ -i\kappa_m \sin \beta_m & \cos \beta_m \end{bmatrix} \quad (11)$$

where $\kappa_m = n_m \sin q_m$ and β_m has been defined above. Once matrices for each individual layer have been calculated, an overall sample matrix M is defined as the product of the individual matrices, so that for a sample with m layers as described in Figure 3 the resultant matrix is defined as:

$$M = \prod_{m=0}^m C_m = \begin{bmatrix} M_{11} & M_{12} \\ M_{21} & M_{22} \end{bmatrix} \quad (12)$$

The reflectivity is then simply related to the matrix elements from M by the relationship:

$$\left| \frac{(M_{11} + M_{12}\kappa_{m+1})\kappa_0 - (M_{21} + M_{22})\kappa_{m+1}}{(M_{11} + M_{12}\kappa_{m+1})\kappa_0 + (M_{21} + M_{22})\kappa_{m+1}} \right|^2 \quad (13)$$

where $m+1$ denotes the substrate and 0 the air. As can be seen this general solution lends itself very well to the use of computers. It is well suited to modelling the reflectivity profiles from samples with complex internal layer structure.

Instrumentation

It is obvious from the definition of the momentum transfer q (Equation 6) that reflectivity measurements as a function of q can be achieved by varying either q or λ . Fixed wavelength reactor sources using monochromatic beams therefore vary q to give the required q range, whereas pulsed neutron sources use a broad band or 'white' beam polychromatic beam and time-of-flight (TOF) to determine λ at fixed q . Common to both methods of measurement however is a high degree of automation which is essential for accurate measurement of $R(q)$. Perhaps the major drawback to neutron reflectivity is the limited number of instruments around the world, which basically only reflects the small number of facilities. However, the number of instruments continues to grow.

This article is not meant to be a comprehensive review of instruments since this has been dealt with in depth in the literature (see for example References 8-17 with a good review by Russell [6]). With two ways of producing neutrons there are basically two designs of reflectometer, each of which will be illustrated by describing typical instruments.

The CRISP reflectometer at the ISIS Facility at the Rutherford Appleton Laboratory is typical of a pulsed source time-of-flight (TOF) instrument. A schematic illustration of the major optical components is shown in Figure 5. The moderated neutrons pass through a double set of choppers, firstly, a disc chopper to select the wavelength range and then a prompt pulse suppressing nimonic chopper. The beam is coarsely collimated by neutron absorbing jaws before entering the experimental measurement area. Fine collimation is achieved by two slits before the sample which define the illuminated area and resolution at the sample position. The beam profile and intensity is monitored just before reaching the sample using a scintillator detector. Post sample the background noise is suppressed by two further slits or by a shielded nose cone. Reflection is detected either with a single or a position sensitive detector. CRISP views a liquid hydrogen moderator which gives a wavelength range of 0.5-6.5 Å, with the disc chopper working at 50 Hz. This range can be extended to 12 Å if the disc chopper is operated at 25 Hz, although this has the drawback of only collecting 1 in every 2 pulses of neutrons. The instrument has been designed to be able to measure liquid surfaces and therefore has a horizontal sample geometry and an inclined beam at 1.5° to the horizontal. At 50 Hz this angle yields a q_z range of 0.05 to 0.65 Å⁻¹, which is optimised for liquid surface measurements. In this q region the kinematic approximation is applicable and is widely utilised for analysis of monolayer structure at the air-water interface [18]. Computer controlled goniometers at the sample position allow movement of solid samples (or confined liquids) to angles other than the 1.5° specified by the instrument design geometry. For liquid surfaces accessing angles less than 1.5° to measure in a q range lower than 0.05 Å⁻¹, a super-mirror is inserted in the path of the beam. This super-mirror totally reflects all wavelengths greater than ~1.5 Å, and reduces the angle of incidence on the sample by where q_{sm} is the angle of the super-mirror to the beam. CRISP like other pulsed neutron source reflectometers has a fixed wavelength range, therefore at each incident angle on the sample a limited q range is obtained. The q range can easily be extended by running 2 or more incident angles and combining the data [9]. The reflectivity profile at a fixed incident angle due to the polychromatic beam is therefore collected all at once, and the time required to measure the total spectrum is only limited by the statistics of the collected data.

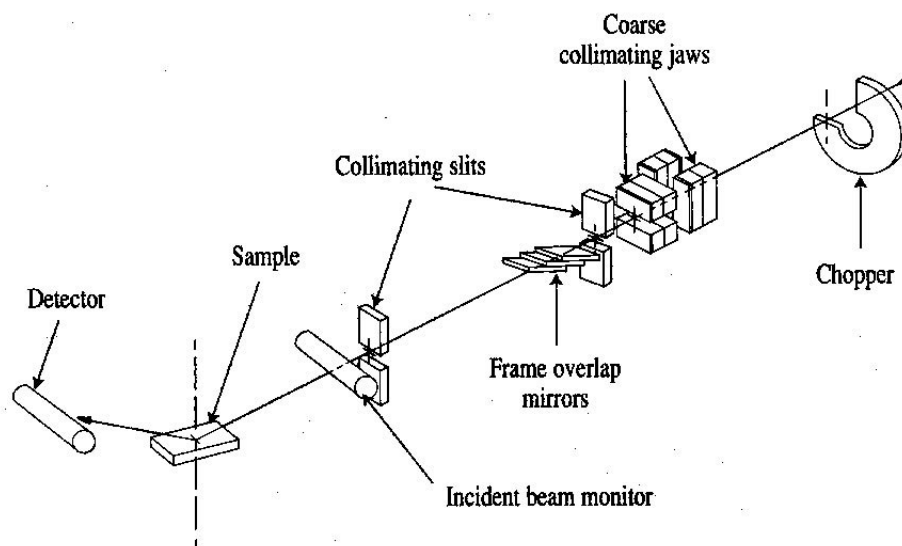


Figure 5: Schematic layout of the time-of-flight neutron reflectometer CRISP at the ISIS Facility, Rutherford Appleton Laboratory.

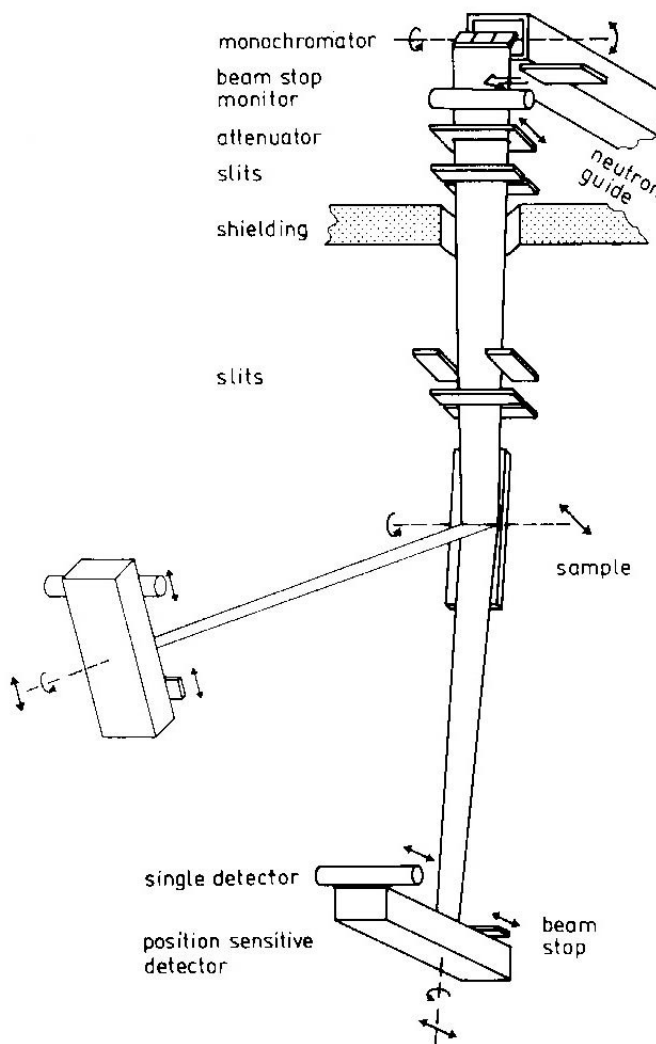


Figure 6: Schematic of the fixed wavelength neutron reflectometer Torema II at the GKSS.

By contrast reactor based reflectometers with fixed wavelength sources measure the reflectivity profile point-by-point in q and hence in $2q$. This method of measurement is often called $q - 2q$ scans and is identical to the method employed by the closely related X-ray reflectometers. A typical example of a $q - 2q$ neutron reflectometer is Torema II at the GKSS research reactor, Hamburg, Germany (see Figure 6). Here moderated neutrons are focused using a monochromating graphite crystal to give a neutron wavelength of 0.43 nm. The beam flux is then monitored before attenuation and 3 variable slits define the beam on the sample. The sample is vertically mounted and is rotated at each angle of measurement (q). The position sensitive detector is rotated at each measurement point at twice the incident angle ($2q$). Due to the relatively high neutron flux at reactor sources, it is necessary at the lower incidence angles to attenuate the beam using plexiglass plates before the collimation to reduce the intensity reaching the sensitive detectors. A drawback of this technique is that both sample and detector must move with respect to the beam. This has a serious repercussion in that the illuminated area varies with rotation of the sample. This is not a problem if there are no lateral inhomogeneities, but can pose difficulties for data interpretation of the measured reflectivity profile if they do exist. On

the whole, many of the advantages or disadvantages regarding sample measurement are trivial and the real differences between instruments now seems to be the perception by the users as to speed of measurement. This is only related to the ‘brightness’ of the neutron source and not the reflectometer.

Isotopic Substitution

• Polymer	Abbreviation	$r_z (\text{\AA}^{-2}) \cdot 10^6$	
		hydrogenated	deuterated
• polystyrene	PS	+1.41	+6.50
• poly(methyl methacrylate)	PMMA	+1.10	+7.22
• polyethylene	PE	-0.33	+8.24
• poly(dimethylsiloxane)	PDMS	+0.06	+5.04
• polybutadiene	PB	+0.41	+6.60

As mentioned before, one of the greatest assets available to neutron reflectivity and of course neutron scattering in general is the use of isotopic substitution for contrast enhancement. For the polymer scientist this is extremely important, because by substituting hydrogen for deuterium a vastly different scattering length density can be produced. This has enormous implications since by a very minor chemical alteration it is possible to contrast a molecule or even a part of a molecule from an environment. In Table 1 the scattering length densities for some common polymers are compared in their normal hydrogenous and deuterated forms using the equation:

$$\rho_z = \frac{\sum b_i D_m N_A}{M_m} \quad (14)$$

where $\sum b_i$ is the sum of the monomer atomic coherent scattering lengths, [4] D_m is the bulk polymer density, N_A is the Avogadro constant and M_m is the monomer molar mass. The use of deuteration can be illustrated further by imagining a reflectivity experiment, for example polymer-polymer interdiffusion, where it is not possible to distinguish between the polymers using X-rays. If two different polymers are chosen their hydrogenous isotopic forms may have enough natural contrast to distinguish between them using neutrons. By comparison of the values in Table 1, however, it is seen that even in this case the differences are very small. Consider now the case of interdiffusion between layers of the same polymer, something which is of vital interest in all aspects of adhesion for example. Here there is no natural contrast and indeed the separate layers are totally identical and no distinction between the sample before and after annealing can be made. If however, one layer is made of deuterated polymer a large difference

is immediately introduced between it and the normal hydrogenous layer. When the layers interdiffuse the change in the interface from a sharp step to a diffuse profile can easily be followed. This deuteration scheme can also be applied to enhance the natural contrast between layers of two different species. Indeed it can be used in a whole host of experiments for example to look at molecules segregating to interfaces and surfaces, or even deuterating the ends of polymer chains to look at preferential adsorption. The use of such deuterated contrast enhancement seem endless and only limited by the ability of the synthetic chemists who provide the labelled polymers.

However, a cautionary note must be added. Although for the most part deuteration leaves a polymer with physically identical properties, for a few well studied examples deuteration can alter considerably the polymer behaviour, especially near phase transitions. Blends of some polymers have been shown to display upper critical solution temperatures contradicting the expected assumption that such isotopic blends form ideal mixtures [19-30]. The effect was first noted for isotopic mixtures of hydrocarbons [31], but has been observed for polystyrene (PS) [19, 22, 23, 26] polybutadiene (PB) [20] and poly(ethylenepropylene) (PEP) [25]. When annealed to equilibrium thin films of isotopic blends of these polymers have a non-homogeneous structure with the lower surface energy component preferentially segregating to the air-polymer and polymer-substrate interfaces. The segregation occurs due to a weakly unfavourable χ interaction parameter originating from the difference in polarizability between C-H and C-D bonds [22]. The effect is only observed however, if the relationship:

$$\chi < \frac{2}{N}$$

(15)

is satisfied, where N is the degree of polymerisation. This means that isotopic segregation in these system is only evident if $N > 2/\chi$ and since χ is typically very small for hydrogenous-deuterated mixtures (χ (h/d-PS) $\gg 2 \times 10^{-4}$ [20]) means that effectively N must be very large. Although this segregation may in principle be a problem in practice it is not so, since careful use of deuteration and design of the experiment can eliminate the problem entirely.

Data Analysis

Routine analysis of reflectivity data would ideally be solved by direct inversion of experimental data into either scattering length density, $r(z)$, or even volume fraction, $f(z)$, profiles. Generally, this cannot be achieved due to the loss of phase information, making this closely related to the phaseless Fourier problem [32]. The analysis then becomes effectively the reverse situation, where reflectivities calculated from simulated $r(z)$ profiles are used in a comparative way to minimise deviations from the measured reflectivity.

For simple systems analytical expressions exist for calculating reflectivity profiles, using the relationships of Fresnel and Snell (see above). However, most real samples do not

typically conform to this simplistic ideal. Extracting the scattering length density profile from the reflectivity data, $R(q)$, becomes necessarily more complex. Because of the lack of phase information from measured reflectivity data it is not generally possible to directly obtain $r(z)$ profiles by inversion of the $R(q)$ curve. Analysis of the $R(q)$ curves to give $r(z)$ profiles is generally not unique in that it is possible that several different $r(z)$ profiles are consistent with the experimental data given the experimental error and limited q range. There are essentially two methods of data analysis:

1. model fitting
2. approximations and free-form techniques.

These are briefly described below together with a review of theoretical work on inversion techniques which are still not widely applicable to real problems due to the limitation with regard to the lack of phase information.

It should be stressed that many of the problems associated with fitting reflectivity data such as uniqueness of fit can be alleviated by use of complementary techniques. In cases where such additional information is not available the careful design of the experiment becomes imperative, and labelling schemes must be extensively used. In general the more information acquired regarding the sample the easier the task of fitting the data.

Model Fitting

The standard approach to reflectivity analysis is model fitting and refinement mostly using the matrix technique. In this method a functional form for $r(z)$ is chosen on the basis of all known data about the system. This will include any complementary information from other reflectivity measurements and/or techniques. The resulting reflectivity profile assuming the $r(z)$ model is computed using Equations 11, 12 & 13 and compared to the experimental data. The parameters of the model are then modified using least-squares or simplex fitting routines so that the deviations of the calculated and measured reflectivity profiles are minimised [33]. Clearly then this approach requires a significant amount of prior knowledge of the sample. This method works well when the system is quite well defined and therefore the initial model $r(z)$ profile gives $R(q)$ sufficiently close to the experimental data. This approach has the advantage that many of the other non-unique solutions for $r(z)$ can be eliminated based on the expected physical properties of the system. However, many cases present themselves where no reasonable starting guess for $r(z)$ exists, making the process of model fitting one of hard work and many intuitive guesses.

All the equations so far used to describe the reflectivity from specimens have been from infinitely sharp interfaces. However, no interface is infinitely sharp, but exhibits a gradient in density not only between the specimen and the surrounding media, i.e. the substrate or air, but also between consecutive layers of polymers. This may represent a diffuse interface between the polymers or even interfacial roughness or waviness. It is important to distinguish between these. Extensive treatment of interfacial roughness and waviness can be found in the literature [34-36]. The difference between roughness and waviness is evident in the $r(z)$ profile projection over the neutron coherent length, l_c [6]. Long-range roughness or waviness of the interface over distances larger than, l_c , has an effect similar to that of a divergence of the incident beam. This can be seen as leading to a small curvature in the $r(z)$ profile. Neutrons impinging on a rough interface, with

interfacial waviness smaller than l_c , have local angles of incident which vary greatly from the average. When projected onto a $r(z)$ profile this gives a wider smooth transition between the two layers. In general, this leads to a reduction in the reflectivity arising from that interface. Névot and Croc [37] showed that for a Gaussian roughness at a bulk interface the Fresnel reflectivity is modified by an exponential factor:

$$R = R_f \exp(-q^2 \sigma^2) \quad (16)$$

where σ is the standard deviation of the Gaussian function describing the roughness. Clearly, from this equation it can be seen that reflectivity is very sensitive to interfacial roughnesses. Large deviations from Fresnel reflectivity are expected even for small amounts of roughness. The treatment of roughness has been extended to thin layers by applying a similar Gaussian factor to the reflectivity coefficients of Equation 7 [38]:

$$r_{j,j+1} = \left(\frac{q_{z,j} - q_{z,j+1}}{q_{z,j} + q_{z,j+1}} \right) \exp\left(-\frac{q^2}{2} \sigma^2\right) \quad (17)$$

An alternative approach to introducing a diffuse interface between two bulk layers is to divide the interface into a number of discrete layers as shown in Figure 7. The change in concentration between each layer must be small enough that the calculated reflectivity profiles do not contain interference fringes associated with the arbitrary choice of histogram thicknesses.

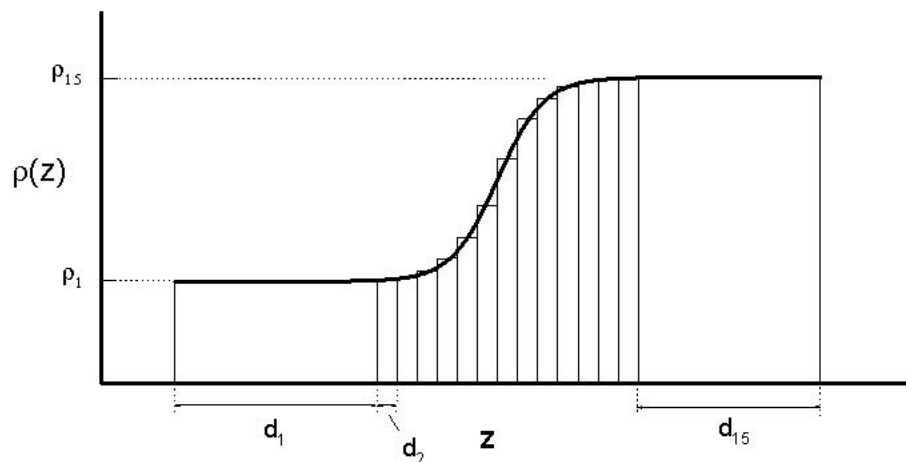


Figure 7: Schematic diagram to show how a diffuse interface, in this case a hyperbolic tangent function, between two polymers can be described by division of the system into histograms with discrete thicknesses, d_i . Associated with each layer is a specific scattering

length density, r_i .

Maximum Entropy and Bayesian Analysis

Since generally the total thickness of the sample is known from other techniques it is possible to apply a free form solution. One approach is the simulated annealing procedure [39], where the total sample thickness is divided into a suitably large number of equally sized pixels. The problem then becomes one of statistically varying the values of r_z in each pixel with certain restrictions concerning jump height and smoothness of curve. This method has the advantage that very little is assumed about the nature of the density profile. However, it does suffer from ambiguity since it is possible to produce a large number of satisfactory fits which are quite possibly meaningless. This problem can be reduced if a suitable starting situation is used. An alternative approach which has been gaining much interest and demonstrated considerable success especially among the polymer community is the maximum entropy (MaxEnt) technique of free form solutions [32, 40, 41]. This uses entropy as a regularising function when varying the value of r_z in each of the pixels, which make up the model. A smoothing correlation function is applied to the values of r_z between pixels removing physically meaningless small wavelength r_z spikes which could occur with uncorrelated pixels. Codes now exist where very little prior knowledge is required other than total sample layer thickness and air and substrate scattering length densities. The program developed by Sivia [42] tries to fit at first one layer and then incrementally more layers to a $r(z)$ model until a precision limit is reached in a χ^2 fit of calculated and experimental $R(q)$. Clearly then these free-form solutions can prove very useful in interpreting data, but a cautionary note must be added that some of the results may still be meaningless.

From the reflectivity data, $R(q)$ it is desired that the scattering length density profile, $r(z)$, can be inferred. This inference can be summarised mathematically by the conditional probability distribution function (PDF), $\text{prob}[r(z) | R(q)]$, [32] where ' $|$ ' means given. This means that with respect to the data, the best estimate of $R(q)$ is given by the $r(z)$ profile which maximises this PDF.

However, this PDF is an unknown, and to calculate it, it is necessary to apply Bayes' Theorem. Bayes' Theorem relates the unknown PDF to two others, one of which can be calculated and the other estimated.

$$\text{prob}[r(z) | R(q)] = \frac{\text{prob}[R(q) | r(z)] \text{prob}[r(z)]}{\text{prob}[R(q)]} \quad (18)$$

Using Bayesian terminology the 'prior' PDF ($\text{prob}[r(z)]$) represents the state of knowledge about $r(z)$ before measurement of the sample. The prior state of knowledge is that modified by the data through a 'likelihood function' ($\text{prob}[R(q) | r(z)]$), which indicates how likely it is that the data set would have been obtained given a trial $r(z)$ profile. To give the required PDF, called the 'posterior' PDF, the product of these 'prior' and 'likelihood' PDF's are taken. This then represents the state of knowledge about the density profile after data have been collected.

It is unusual to be totally ignorant of prior knowledge about the sample, since information is usually available on the sample preparation, in addition to the possibility of being able to describe the $r(z)$ profile by a histogram-like structure of 'heights' and 'widths'. This additional prior knowledge allows the problem to be simplified further, and an estimate of the $r(z)$ profile is then obtained by maximising the 'posterior' PDF, $\text{prob}[\{d_i, r_i\} | R(q), r_s]$,

where d_i and r_i are thicknesses and scattering length densities of each of the M layers describing $r(z)$, and r_s is the substrate scattering length density. If the number of layers is known this is then a simple problem of parameter estimation. If, however, M is not known it is necessary to calculate the PDF $\text{prob}[M \frac{1}{2}R(q), r_s]$ [32, 40].

Speckle Holography and Other Techniques

Although the application of neutrons to the study of density profiles as a function of depth is now well established, the information obtained is often subject to questions of uniqueness. In fact it is quite possible that several different models adequately describe the same reflectivity profile. The experimental data hold vital information about layer thicknesses as well as interfacial and surface structure. As described above most work to date has involved fitting parameterised models to the measured data. This is mostly successful, but can lead to the possibility of scientific prejudice leading to constraints of the model. To alleviate this problem, the ability to directly invert the reflectivity data to a density-depth profile would be most desirable. However, the lack of phase information in the reflectivity data prevents true inversion, and while several inversion methods can be applied they still rely on some knowledge of phase information. This is discussed in more detail in section 6.4. Two related approaches to reduce ambiguity of fits, speckle holography and use of magnetic substrates, are briefly described below.

To alleviate the ambiguity problems inherent in traditional reflectivity measurements, a novel approach is based on the holographic speckle imaging technique [43, 44] used in optical astronomy, which Sivia [32, 45] has called speckle holography. Assuming that the reflectivity is related to the derivative of the density profile (dr/dz) by a phaseless Fourier transform:

$$R(q) = \frac{(4\pi)^2}{q^4} \left| \int_{-\infty}^{+\infty} \frac{d\rho}{dz} e^{iqz} dz \right|^2 \quad (19)$$

then the autocorrelation function (ACF) of dr/dz is given by the Fourier transform of $R(q)q^4$:

$$ACF\left(\frac{d\rho}{dz}\right) = c \int_{-\infty}^{+\infty} R(q)q^4 e^{-iqz} dq \quad (20)$$

where c is a constant. However, Sivia shows [32, 45] that autocorrelation functions are not unique unless there is a distant reference signal (see Figure 8). Such a reference signal can be introduced into a reflectivity sample by introducing a thick buffer layer between the real sample under investigation and the substrate. The $r(z)$ discontinuity at the substrate-buffer layer interface then acts as the distant reference in the dr/dz plot. In practice, the reflectivity data plotted as Rq^4 versus q is Fourier transformed to give the autocorrelation function of dr/dz . The $r(z)$ profile is then simply obtained by integration of the peaks in the $ACF(dr/dz)$ profile. Although the Fourier approximation of Equation 19 is only strictly valid for $R(q) \ll 1$ and $q \gg q_c$, this method at least in principle should distinguish between previously ambiguous solutions.

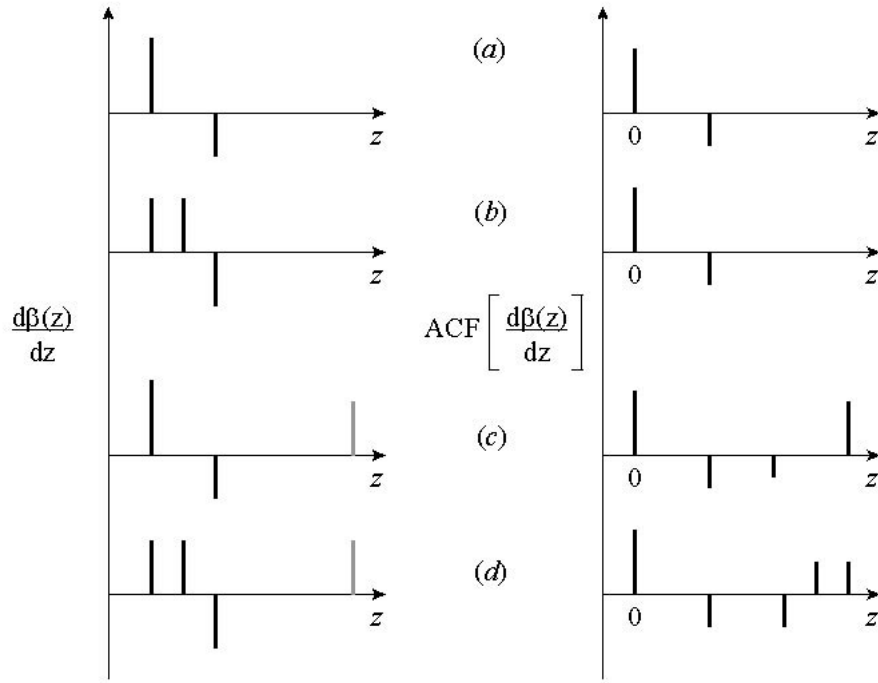


Figure 8: Derivatives of the scattering length density profiles ($d\beta/dz$) from 4 different samples, together with their autocorrelation functions ($ACF(d\beta/dz)$). The values of $d\beta/dz$ in (a) and (c) and (b) and (d) are identical except for the inclusion of a reference point source on the right-hand side of (c) and (d). The inclusion of the reference source produces a scaled mirrored copy of the $d\beta/dz$ on the right-hand side of the ACF's (After References [32] and [45]).

Direct inversion of simulated data has shown that simultaneous use of X-ray and neutron reflectivity data on a sample with an additional artificial bilayer between the substrate and the film under investigation [46], can often eliminate ambiguities in data inversion [32]. This idea has been developed by Sanyal [47] for use with tuneable X-rays and the distorted-wave Born Approximation to successfully analyse synchrotron reflectivity data obtained from Langmuir-Blodgett films. The neutron equivalent has also been developed and is based on the use of polarised neutrons and magnetised substrates, to determine uniquely and unambiguously the density profile of thin non-magnetic films [48].

The essential requirement of this technique is to be able to measure the unknown non-magnetic film on two different substrates whose scattering-length densities can be denoted by r_1 and r_2 . Using a magnetised substrate and polarised neutrons this condition is satisfied for the scattering-length densities of the parallel and anti-parallel spin components of the polarised neutrons. In this way two simultaneous experiments are being performed on the same film but effectively different substrates. The scattering-length densities of the magnetic substrate will then be given by either the sum or difference of the nuclear and magnetic components. Under these conditions the difference in the reflectivities for the two neutron spin states is given by:

$$\frac{(R_1(q) - R_2(q))}{R(q)} = (\rho_1^2 - \rho_2^2) + 2(\rho_1 - \rho_2) \text{Re} \left[\int \frac{d\rho(z)}{dz} e^{iqz} dz \right] \quad (21)$$

Where z is the perpendicular distance to the surface, $R_i(q)$ are the reflectivities as a function of the neutron wavevector transfer, q , of the two spin states of the neutrons ($i=1,2$) and Fresnel reflectivity ($i=f$) and q' is the average wavevector transfer measured in the film. In the simplest approximation $R_f(q)$ is given by $4\pi/q^4$. If Equation (21) is Fourier transformed, the derivative of the scattering length density, $dr(z)/dz$, of the film is obtained directly and unambiguously:

$$\begin{aligned} & \frac{1}{2\pi} \int \left(\frac{R_1(q) - R_2(q)}{R_f(q)} \right) e^{-iq'z} dz \\ &= (\rho_1 - \rho_2) \delta(0) + (\rho_1 - \rho_2) \left[\frac{d\rho(z)}{dz} + \frac{d\rho(-z)}{dz} \right] \end{aligned} \quad (22)$$

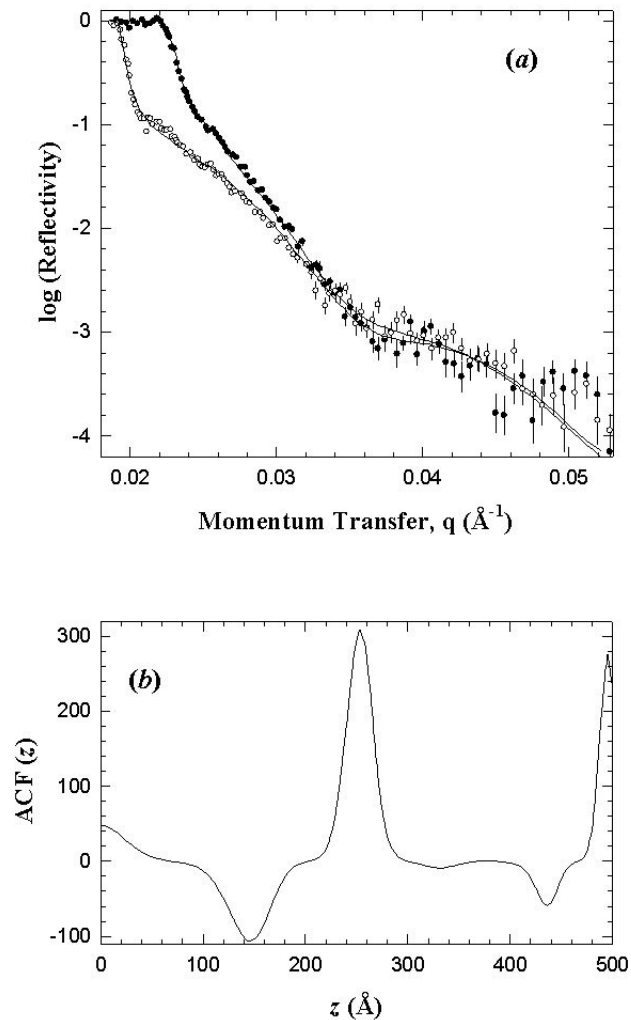


Figure 9: (a) Reflectivity curves of a dPS film on a nickel substrate measured using polarised neutrons with spins parallel (solid circles) and anti-parallel (open circles) to the magnetic field. (b) Autocorrelation function (ACF) data obtained by applying Equation 22 to the data of (a). The spike in the data at a $z=25$ nm represents the thickness of the film.

The delta function, $d(0)$, arises from the interface between a substrate and the film deposited on it. Using polarised neutrons the reflectivities of $R_1(q)$ and $R_2(q)$ are obtained from the two spin states of the neutron when the sample is in a magnetic field. This technique is ideally suited to the study of polymer films, and preliminary experiments to demonstrate its effectiveness have been successfully demonstrated using a nickel substrate to determine the profile of a single dPS film, see Figure 9 [49]. Of course the technique is not restricted to single layers as demonstrated by the simulations of Sivia [48]. This study showed, however, that a perfectly sharp interface between the magnetic substrate and the non-magnetic film under investigation is vital for the distorted-wave Born Approximation to be valid.

Inversion Techniques

The ideal situation for analysis of reflectivity data would be a direct inversion of the experimental data into a $r(z)$ profile. However, a big obstacle to this becoming routine is the loss of phase information incurred due to the square term in the reflectivity coefficients $R(q) = \frac{1}{2} r_{01/2}^2$. Due to this, there are presently no valid analytic expressions relating $R(q)$ to $r(z)$ over the entire q_z range of the reflectivity curve. However, in a limited number of cases direct inversion is possible. The kinematic approximation for example together with extensive deuteration schemes is very powerful for systems within the kinematic regime, i.e. in the limit of high q ($q \gg q_c$) and which are weakly reflecting ($R \gg 1$). This limits the approach to systems which form very thin layers at bulk interfaces. The technique is well established therefore for the interpretation of reflectivity data of surfactants at the air-water interface (see for example references 18, 50, 51). The kinematic approximation has been applied to polymers adsorbed at the air-water interface [52] but is of limited use to most polymer systems.

True inversion as stated requires knowledge of not only the amplitude but also phase of the reflection coefficients, r_{01} . If these were known then in principle inversion of $R(q)$ into $r(z)$ profiles would be possible and practical algorithms exist. [53, 54]. However, in general, except in the cases discussed above, no phase information exists and methods to retrieve it from the measured reflectivity have been discussed on model simulated data in the literature [55-66]. However, many of these methods can yield numerous solutions for the phase, which then leads to non-unique inversion solutions. Extra data are then required to identify the correct solution, either by complementary depth profiling techniques [67, 68], or isotopic substitution methods [50, 51]. As seen above when placed on a magnetic reference substrate the reflectivity data from an unknown film which is non-magnetic can be inverted to $r(z)$ with the use of polarised neutrons. Schemes for inverting the two spin states of the reflectivity data have been proposed by Sivia [48] and Majkrzak [69]. However these methods are based on the Born approximation which is valid only in the kinematic limit ($q \gg q_c$). Independently de Haan [70] and Majkrzak [65, 71] both proposed essentially identical methods for retrieving exactly the amplitude and phase information of the reflection coefficient even in the dynamic regime ($q \gg q_c$). This method uses a 'tuneable' reference layer beneath the unknown layer. The complex reflection coefficient of the unknown layer can be obtained unambiguously by measuring the sample with 3 spin states with respect to the magnetic reference layer [66].

References

1. Lekner J, Theory of Reflection , 1987, Martinus Nijhoff, Dordrecht
2. Higgins JS & Benoit HC, Polymers and Neutron Scattering , 1994, Clarendon Press, Oxford
3. Koester L, in Bonse U & Rauch H (eds), Neutron Interferometry , 1979, Oxford Science Publications, Oxford
4. Sear VF, Neutron News , 1992 3, 26
5. Born M & Wolf E, Principles of Optics , 1980, Pergamon Press, Oxford, 6 th edition
6. Russell TP, Mater Sci Rep , 1990, 5, 171
7. Heavens OS, Optical Properties of Thin Films , 1955, Butterworth, London
8. Penfold J, Physica B , 1991, 173 , 1
9. Bucknall DG, Penfold J, Webster JRP, et al , ICANS XIII Proceedings, PSI Proc 95-02 , 1995, 1, 123
10. Stamm M, Hüttenbach S & Reiter G, Physica B , 1991, 173 , 11
11. Karim A, Arendt BH, Goyette R, Huang YY, Kleb R & Felcher GP, Physica B , 1991, 173 , 17
12. van Well AA, de Haan VO & Fredrikze H, Physica B , 1994, 198 , 217
13. Clemens D, Physica B , 1996, 221 , 507
14. Sun X, Bouchard E, Lapp A, Farnoux A, Daoud M and Jannick G, Europhys Letts , 1988, 6, 207
15. Sun X, Farnoux B, des Cloizeaux J & Jannick G, Physica B , 1989, 156/157 , 407
16. Anastasiadis SH, Russell TP, Satija SK & Majkrzak CF, J Chem Phys , 1990, 92 , 5677
17. Vaknin D, Kjaer K, Als-Nielsen J & Lösche M, Makromol Chem Macromol Symp , 1991, 46 , 383
18. Lu JR, Hromadova M, Simister EA, Thomas RK & Penfold J, J Phys Chem , 1994, 98 , 11519
19. Bates FS, Wignall GD & Koehler WC, Phys Rev Lett , 1985, 55 , 2425
20. Bates FS & Wignall GD, Phys Rev Lett , 1986, 57 , 1429
21. Pan DH & Prest Jr WM, J Appl Phys , 1985, 58 , 15
22. Jones RAL, Kramer EJ, Rafailovich MH, Sokolov J & Schwarz SA, Phys Rev Lett , 1989, 62 , 280
23. Jones RAL & Kramer EJ, Philosophical Mag B , 1990, 62 , 129

24. Leonhardt DC, Johnson HE & Granick S, *Macromolecules* , 1990, 23 , 685
25. Jones RAL, Norton LJ, Kramer EJ, Bates FS & Wiltzius P, *Phys Rev Lett* , 1991, 66 , 1326 & Jones RAL, *Phys Rev Lett* , 1991, 66 , 3087
26. Zhao W, Zhao X, Rafailovich MH, Sokolov J, Mansfield T, Stien RS, Composto RC, Kramer EJ, Jones RAL, Sansome M & Nelson M, *Physica B* , 1991, 173 , 43
27. Hariharan A, Kumar SK & Russell TP, *J Chem Phys* , 1993, 98 , 4163
28. Hariharan A, Kumar SK, Rafailovich MH, Sokolov J, Zheng X, Duong D, Schwarz SA & Russell TP, *J Chem Phys* , 1993, 99 , 656
29. Graessley WW, Krishnamoorti R, Balsara NP, Fetters LJ, Lohse DJ, Schulz DN & Sissano JA, *Macromolecules* , 1993, 26 , 1137
30. Hong PP, Boerio FJ & Smith SD, *Macromolecules* , 1994, 27 , 596
31. Bartell LS & Roskos RR, *J Chem Phys* , 1966, 44 , 457
32. Sivia DS, Hamilton WA & Smith GS, *Physica B* , 1991, 173 , 121
33. Press WH, Teukolsky SA, Vetterling WT & Flannery, *Numerical Recipes* , 2 nd ed, Cambridge University Press, Cambridge, 1992
34. Sinha SK, Sirota EB, Garoff S & Stanley HB, *Phys Rev B* , 1988, 38 , 2297
35. Pynn R, *Phys Rev B* , 1992, 45 , 602
36. Pynn R & Baker S, *Physica B* , 1994, 198 , 1
37. Névoit L & Crocé P, *Phys Appl* , 1980, 15 , 761
38. Cowley RA & Ryan TW, *J Phys D* , 1987, 20 , 61
39. Kunz K, Reiter J, Götzelman A, Reiter J & Stamm M, *Macromolecules* , 1993, 26 , 4316
40. Geoghegan M, Jones RAL, Sivia DS, Penfold J & Clough AS, *Phys Rev E* , 1996, 53 , 825
41. Sivia DS, *Data Analysis - A Bayesian Tutorial* , 1996, Clarendon Press, Oxford
42. Sivia DS, Private communication
43. Weigert G & Baier G, *Astron Astrophys* , 1985, 150 , L18
44. Labeyrie A, *Astron Astrophys* , 1970, 6, 85
45. Sivia DS, Hamilton WA, Smith GS, Rieker TP & Pynn R, *J Appl Phys* , 1991, 70 , 732
46. Lesslauer W & Blasie JK, *Acta Cryst* , 1971, A27 , 456
47. Sanyal MK, Sinha SK, Gibaud A, Huang KC, Calvalho BL, Rafailovich M, Sokolov J, Zhao X & Zhao W, *Europhys Letts* , 1993, 21 , 691

48. Sivia DS & Pynn R, SPIE Conference Proc, Neutron Optical Devices and Applications , 1992, 1738 , 305
49. Penfold J, Webster JRP, Bucknall DG & Sivia DS, Colloids and Surfaces A: Physicochem Eng Aspects , 1994, 86 , 165
50. Lu JR, Li ZX, Smallwood J, Thomas RK & Penfold J, J Phys Chem , 1995, 99 , 8233
51. Crowley TL, Lee EM, Simister EA & Thomas RK, Physica B , 1991, 173 , 143
52. Henderson JA, Richards RW, Penfold J & Thomas RK, Macromolecules , 1993, 26 , 65
53. Moses HE, Phys Rev , 1956, 102 , 559
54. Roberts TM, Physica B , 1991, 173 , 143
55. Burge RE, Fiddy MA, Greenaway AH & Ross G, Proc Roy Soc London , 1976, A350 , 191
56. Klibanov MV & Sacks PE, J Math Phys , 1992 33 , 3813
57. Klibanov MV & Sacks PE, J Comput Phys , 1994, 112 , 273
58. Lipperheide R, Reiss G, Fiedeldey H, Sofianos SA & Leeb H, Physica B , 1993, 190 , 377
59. Reiss G & Lipperheide R, Phys Lett A , 1994, 196 , 133
60. Lipperheide R, Reiss G, Leeb H, Fiedeldey H & Sofianos SA, Phys Rev B , 1995, 51 , 11032
61. Lipperheide R, Fiedeldey H, Leeb H, Reiss G & Sofianos SA, Physica B , 1995, 213&214 , 914
62. Reiss G & Lipperheide R, Phys Rev B , 1996, 53 , 8157
63. Reiss G, Physica B , 1996, 221 , 533
64. Lipperheide R, Reiss G, Leeb H & Sofianos SA, Physica B , 1996, 221 , 514
65. Majkrzak CF & Berk NF, Physica B , 1996, 221 , 520
66. de Haan VO, van Well AA, Sacks PE, Adenwalla S & Flecher GP, Physcia B , 1996, 221 , 524